

[illegible]

Claims 1-12 have been cancelled. New claims 13-21 have been added further to define the invention, over U.S. Application Serial No. 09/020,018 of which the application filed herewith is a divisional application.

Respectfully submitted,

By

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MARKED UP AMENDED SPECIFICATION PARAGRAPHS

Page 9, first complete paragraph

The plasmids encoding SakSTAR(K35A,E38A,K74A,E75A), SakSTAR(E38A,E75A,R77A), SakSTAR(E38A,E75A), SakSTAR(K35A,E75A,R77A), SakSTAR(K35A,E75A), SakSTAR(E80A), SakSTAR(D82A), SakSTAR(E75A,D82A), SakSTAR(K74A) and SakSTAR(E75A) were constructed by the spliced overlap extension polymerase chain reaction (SOE-PCR) (24), using Vent DNA polymerase (New England Biolabs, Leusden, The Netherlands), and available or generated *sakSTAR* variants as template. Two fragments were amplified by PCR, the first one starting from the 5' end of the *staphylokinase* gene with primer 5'-CAGGAAACAGAATTCAGGAG-3' (SEQ ID NO: 1) to the region to be mutagenized (forward primer), the second one from the same region (backward primer) to the 3' end of the *staphylokinase* gene with primer 5'-CAAAACAGCCAAGCTTCATTCATTCAGC-3' (SEQ ID NO: 2).

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The variants SakSTAR(Y17A,F18A), SakSTAR(F104A), SakSTAR(F111A), SakSTAR(Y9A), SakSTAR(Y91A), SakSTAR(Y92A), SakSTAR(I87A), SakSTAR(I106A) and SakSTAR(I120A) were constructed with the Chameleon Double-Stranded Site-Directed Mutagenesis kit from Stratagene (La Jolla, USA), using the *pMEX.SakSTAR* vector as template, and following instructions of the supplier. The mutagenic oligonucleotides (not shown) were used in combination with the selection-primer LY34 5' CAAAACAGCCGAGCTTCATTCATTCAGC (SEQ ID NO: 3), which destroys the unique *HindIII* site located 3' to the *staphylokinase* encoding gene in *pMEX.SakSTAR* and allows to counter-select the non-mutant progeny by *HindIII* digestion. The deletion of the *HindIII* site was in most cases correlated with the presence of the desired mutation introduced by the

mutagenic oligonucleotide. The variant SakSTAR(I133A), was constructed by performing a polymerase chain reaction on the *pMEX.SakSTAR* plasmid using the primer 818A located at the 5' end of the *sakSTAR* gene (5' CAGGAAACAGAATTCAGGAG) (SEQ ID NO: 1) and the mutagenic primer LY58 (5' TTCAGCATGCTGCAGTTATTTCTTTTCTGCAACAACCTTGG) (SEQ ID NO: 4). The amplified product (30 cycles: 30 sec at 94°C, 30 sec at 50°C, 30 sec at 72°C) was purified, digested with EcoRI and PstI, and ligated into the corresponding sites of *pMEXSakSTAR*.

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The plasmids encoding all the other variants listed in Table 3 were constructed by direct PCR or by the spliced overlap extension polymerase chain reaction (SOE-PCR)(24) using *pMEX.SakSTAR* or available plasmids encoding SakSTAR variants as template. Two fragments were amplified by PCR (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C), the first one starting from the 5' end (primer 818A) of the *staphylokinase* gene to the region to be mutagenized (forward primer), the second one from this same region (backward primer) to the 3' end of the gene with primer 818D (5' CAAACAGCCAAGCTTCATTCATTCAGC) (SEQ ID NO: 5). The forward and backward primers shared an overlap of around 24 bp (primers not shown). The two purified fragments were then assembled together in a second PCR reaction with the external primers 818A and 818D (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C). The amplified product from this final reaction was purified, digested with EcoRI and HindII and ligated into the corresponding site of *pMEX.SakSTAR*. For each construction, the sequence of the variant was confirmed by sequencing the entire *SakSTAR* coding region.

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The variants SakSTAR(K102C) and SakSTAR(K109C), were constructed by the spliced overlap extension polymerase chain reaction (SOE-PCR) (24) using *pMEX.SakSTAR* encoding SakSTAR as template. Two fragments were amplified by PCR (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C), the first one starting from the 5' end (primer 818A) of the *staphylokinase* gene to the region to be mutagenized (forward primer), the second one from this same region (backward primer) to the 3' end of the gene with primer 818D (5' CAAACAGCCAAGCTTCATTCATTCAGC) (SEQ ID NO: 5). The forward and backward primers shared an overlap of around 24 bp (for the construction of K102C: TAT GAT AAG AAT TGC AAA AAA GAA GAA (backward) (SEQ ID NO: 6) and TTC TTC TTT TTT GCA ATT CTT ATC ATA (forward) (SEQ ID NO: 7) for the construction of K109C: AAA AAG AAG AAA CGT GCT CTT TCC CTA (backward) (SEQ ID NO: 8) and TAG GGA AAG AGC ACG TTT CTT CTT TTT [(forward)] (forward) (SEQ ID NO: 9). The two purified fragments were then assembled together in a second PCR reaction with the external primers 818A and 818D (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C). The amplified product from this final reaction was purified, digested with EcoRI and HindIII and ligated into the corresponding site of *pMEX.SakSTAR*. For each construction, the sequence of the variant was confirmed by sequencing the entire *SakSTAR* coding region.

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CIP of US 08/371,505 (Jan 11, 1995)

CIP of US 08/499,092 (July 6, 1995)

CIP of US 08/784,971 (Jan 16, 1997)

This invention relates to new staphylokinase derivatives with reduced immunogenicity, their identification, production and use in the treatment of arterial thrombosis and the preparation of a pharmaceutical composition for treating arterial thrombosis. More in particular it relates to the use of engineered staphylokinase derivatives for the preparation of a pharmaceutical composition for treating myocardial infarction.

Staphylokinase, a protein produced by certain strains of *Staphylococcus aureus*, which was shown to have profibrinolytic properties more than 4 decades ago (1,2) appears to constitute a potent thrombolytic agent in patients with acute myocardial infarction (3,4). The staphylokinase gene has been cloned from the bacteriophages *sak*ϕC (5) and *sak*42D (6) as well as from the genomic DNA (*sak*STAR) of a lysogenic *Staphylococcus aureus* strain (7). The staphylokinase gene encodes a protein of 163 amino acids, with amino acid 28 corresponding to the NH₂-terminal residue of full length mature staphylokinase (6,8,9). The mature protein sequence of the wild-type variant SakSTAR (9) is represented in Figure 1. Only four nucleotide differences were found in the coding regions of the *sak*ϕC, *sak*42D and *sak*STAR genes, one of which constituted a silent mutation (6,8,9).

In a plasma milieu, staphylokinase is able to dissolve fibrin clots without associated fibrinogen degradation (10-12). This fibrin-specificity of staphylokinase is the result of reduced inhibition by α_2 -antiplasmin of plasmin.staphylokinase complex bound to fibrin, recycling of staphylokinase from the plasmin.staphylokinase complex following inhibition by α_2 -antiplasmin, and prevention of the conversion of circulating plasminogen.staphylokinase to plasmin.staphylokinase by α_2 -antiplasmin (13-15). In addition staphylokinase has a weak affinity for circulating but a high affinity for fibrin-bound plasminogen (16) and staphylokinase requires NH_2 -terminal processing by plasmin to display its plasminogen activating potential (17). In several experimental animal models, staphylokinase appears to be equipotent to streptokinase for the dissolution of whole blood or plasma clots, but significantly more potent for the dissolution of platelet-rich or retracted thrombi (18,19).

Staphylokinase is a heterologous protein and is immunogenic in man. The intrinsic immunogenicity of staphylokinase, like that of streptokinase, clearly hampers its unrestricted use. Not only will patients with preexisting high antibody titers be refractory to the thrombolytic effect of these agents, but allergic side effects and occasional life-threatening anaphylaxis may occur (20). Because both streptokinase and staphylokinase are heterologous proteins, it is not obvious that their immunogenicity could be reduced by protein engineering. Indeed, no successful attempts to generate active low molecular weight fragments from streptokinase have been reported. In staphylokinase, deletion of the NH₂-terminal 17 amino acids or the COOH-terminal 2 amino acids inactivates the molecule, which in addition is very sensitive to inactivation by site-specific mutagenesis (21).

Nevertheless, we have, surprisingly, found that the wild-type staphylokinase variant SakSTAR (9) contains three non-overlapping immunodominant epitopes, at least two of which can be eliminated by specific site-directed mutagenesis, without inactivation of the molecule (22). These engineered staphylokinase variants are less reactive with antibodies elicited in patients treated with wild-type staphylokinase, and are significantly less immunogenic than wild-type staphylokinase, as demonstrated in rabbit and baboon models and in patients with peripheral arterial occlusion (22).

The present invention relates to general methods for the identification, production and use of staphylokinase derivatives showing a reduced antigenicity and immunogenicity as compared to wild-type staphylokinase. The derivatives have essentially the amino acid sequence of wild-type staphylokinase or modified versions thereof and essentially intact biological activities, but have a reduced reactivity with a panel of murine monoclonal antibodies and/or with antibodies induced in patients by treatment with wild-type SakSTAR.

The invention also relates to a method for producing the derivatives of the invention by preparing a DNA fragment comprising at least the part of the coding sequence of staphylokinase that provides for its biological activity; performing *in vitro* site-directed mutagenesis on the DNA fragment to replace one or more codons for wild-type amino acids by a codon for another amino acid; cloning the mutated DNA fragment in a suitable vector; transforming or transfecting a suitable host cell with the vector; culturing the host cell under

conditions suitable for expressing the DNA fragment and purifying the expressed staphylokinase derivative to homogeneity; preferably the DNA fragment is a 453 bp *EcoRI-HindIII* fragment of the plasmid *pMEX602sakB* (22,23), the *in vitro* site-directed mutagenesis is preferably performed by spliced overlap extension polymerase chain reaction with Vent DNA polymerase (New England Biolabs) or Taq polymerase (Boehringer Mannheim) and with available or generated wildtype *sakSTAR* or *sakSTAR* variants as template (24).

The invention also relates to pharmaceutical compositions comprising at least one of the staphylokinase derivatives according to the invention together with a suitable excipient, for treatment of arterial thrombosis. Pharmaceutical compositions, containing less immunogenic staphylokinase variants as the active ingredient, for treating arterial thrombosis in human or veterinary practice may take the form of powders or solutions and may be used for intravenous, intraarterial or parenteral administration. Such compositions may be prepared by combining (e.g. mixing, dissolving etc.) the active compound with pharmaceutically acceptable excipients of neutral character (such as aqueous or non-aqueous solvents, stabilizers, emulsifiers, detergents, additives), and further, if necessary with dyes.

Furthermore the invention relates to the use of the staphylokinase derivatives for the treatment of arterial thrombosis, in particular myocardial infarction, and to the use of staphylokinase derivatives for the preparation of a pharmaceutical composition for the treatment of arterial thrombosis, in particular myocardial infarction.

In the above and the following the terms "derivatives", "mutants" and "variants" are used interchangeably.

The present invention will be demonstrated in more detail in the following examples, that are however not intended to be limiting to the scope of the invention. Based on the present invention other variants and improvements will be obvious for the person skilled in the art. Thus random mutagenesis is likely to generate alternative mutants with reduced immunogenicity and possibly increased functional activity, whereas deletions or substitution with other amino acids may yield additional variants with reduced immunogenicity.

EXAMPLE 1

Epitope mapping of wild-type staphylokinase

✕ The epitope specificity of a panel of 15 murine MAbs (22) raised against wild-type SakSTAR was determined by real-time biospecific interaction analysis (BIA) with the BIAcore instrument (Pharmacia, Biosensor AB, Uppsala, Sweden). The MAbs were immobilized on the surface of the Sencor Chip CM5 with the Amine Coupling Kit (Pharmacia Biosensor AB) as recommended by the manufacturer (25). Immobilization was performed from protein solutions at a concentration of 20 µg/mL in 10 mmol/L sodium acetate at pH 5.0 at a flow rate of 5 µL/min during 6 minutes. This resulted in covalent attachment of 5,000 to 10,000 resonance unit (RU) of antibody (corresponding to 0.035 to 0.07 pmol/mm²). The SakSTAR solutions were passed by continuous flow at 20°C past the sensor surface. At least four concentrations of each analyte (range, 50 nmol/L to 50 µmol/L) in 10 mmol/L HEPES, 3.4 mmol/L EDTA, 0.15 mol/L NaCl, and 0.005% Surfactant P20, pH 7.2, were injected at a flow rate of 5 µL/min during 6 minutes in the association phase. Then sample was replaced by buffer, also at a flow rate of 5 µL/min during 6 minutes. After each cycle, the surface of the sensor chip was regenerated by injection of 5 µL of 15 mmol/L HCl. Apparent association (k_{ass}) and apparent dissociation (k_{diss}) rate constants were derived from the sensorgrams as described in detail elsewhere (26), and association equilibrium constants (K_A) calculated as their ratio.

Determination of the equilibrium association constants for the binding of wild-type and variant SakSTAR to insolubilized MAbs (Table 1) yielded apparent association constants of 10^7 to 10^8 (mol/L)⁻¹, which are one to two orders of magnitude lower than the apparent association constants previously obtained for the binding of these MAbs to insolubilized wild-type SakSTAR (22). ✕ If the MAbs instead of the SakSTAR variants are insolubilized, avidity effects of the bivalent MAbs are indeed avoided. The present values are indeed in better agreement with known association constants of Mabs, and therefore this “reversed” procedure was used throughout the present invention.

In the tables the column indicated with "Variant" states the various staphylokinase derivatives which are identified by listing between brackets the substituted amino acids in single letter symbols followed by their position number in the mature staphylokinase sequence and by the substituting amino acids in single letter symbol; the column "Exp." indicates expression levels in mg/L, and the column "Spec. Act." indicates the specific activity in Home Units as defined in example 2. Indications "17G11", "26A2" etc. refer to monoclonal antibodies binding to the indicated epitopes I, II and III (22). Epitope I is recognized by the antibody cluster 17G11, 26A2, 30A2, 2B12 and 3G10, whereas epitope II is recognized by the antibody cluster 18F12, 14H5, 28H4, 32B2 and 7F10, and epitope III by the antibody cluster 7H11, 25E1, 40C8, 24C4 and 1A10. Human plasma "Pool" indicates a plasma pool from initially 16 and subsequently 10 patients immunized by treatment with SakSTAR, "Subpool B" indicates a plasma pool from three patients that absorbed less than 50% of the induced antibodies with SakSTAR(K35A,E38A,K74A,E75A,R77A) and "Subpool C" indicates a plasma pool from 3 patients that absorbed >90% of the induced antibodies with SakSTAR(K35A,E38A,K74A,E75A,R77A) (22). In tables 6, 7 and 8 an additional pool of plasma from 40 patients immunized by treatment with SakSTAR (Pool 40) was also used.

EXAMPLE 2

Construction, epitope mapping with murine monoclonal antibodies and absorption with pooled plasma of immunized patients, of "alanine-to-wild-type" reversal variants of "charged-cluster-to-alanine" mutants of staphylokinase

As stated above, wild-type staphylokinase (SakSTAR variant (9)) contains three non-overlapping immunodominant epitopes, two of which can be eliminated by specific site-directed substitution of clusters of two (K35A,E38A or E80A,D82A) or three (K74A,E75A,R77A) charged amino acids with Ala (22). The combination mutants SakSTAR(K35A,E38A,K74A,E75A,R77A) in which Lys35, Glu38, Lys74, Glu75 and Arg77, and SakSTAR(K74A,E75A,R77A,E80A,D82A) in which Lys74, Glu75, Arg77, Glu80 and Asp82 were substituted with Ala (previously identified as SakSTAR.M3.8 and

SakSTAR.M8.9, respectively (22)), were found to have a reduced reactivity with murine monoclonal antibodies against two of the three immunodominant epitopes and to absorb on average only 2/3 of the neutralizing antibodies elicited in 16 patients by treatment with wild-type SakSTAR (22). These mutants also induced less antibody formation than wild-type SakSTAR in experimental thrombolysis models in rabbits and baboons, and in patients with peripheral arterial occlusion (22). However, their specific activities were reduced to approximately 50% of that of wild-type SakSTAR, which would be of some concern with respect to the clinical use of these compounds.

In an effort to improve the activity and stability without loss of the reduced antibody recognition, the effect of a systematic reversal of one or more of these substituted amino acids to the wild-type residues was studied. Fourteen new mutants were constructed, purified and characterized in terms of specific activity, reactivity with the panel of murine monoclonal antibodies, and absorption of antibodies from plasma of patients treated with wild-type SakSTAR (Table 1). The present example thus focusses on reversal from alanine to the wild-type residue of one or more of the seven amino acids of SakSTAR listed above i.e. K35, E38, K74, E75, R77, E80 and D82.

Reagents and Methods

The source of all reagents used in the present study has previously been reported (22). Restriction enzymes were purchased from Pharmacia (Uppsala, Sweden) or Boehringer Mannheim (Mannheim, Germany). T4 DNA ligase, Klenow Fragment of *E. coli* DNA polymerase I and alkaline phosphatase were obtained from Boehringer Mannheim. Enzyme reactions were performed using the conditions suggested by the suppliers. Plasmid DNA was isolated using a QIAGEN-purification protocol (provided by Westburg, Leusden, The Netherlands). *pMEX.602sakB* (i.e. *pMEX.SakSTAR*) was constructed as described elsewhere (23). SakSTAR, SakSTAR(K35A,E38A), SakSTAR(K74A,E75A,R77A), SakSTAR(E80A,D82A), SakSTAR(K35A,E38A,K74A,E75A,R77A) and SakSTAR(K74A,E75A,R77A,E80A,D82A) were produced and purified as described elsewhere (22). Transformations of *E. coli* were performed utilizing the calcium phosphate

procedure. DNA sequencing was performed using the dideoxy chain termination reaction method and the Automated Laser fluorescent A.L.F.TM (Pharmacia). The chromogenic substrate (S2403) L-Pyroglutamyl-L-phenylalanyl-L-lysine-p-nitroaniline hydrochloride was purchased from Chromogenix (Belgium). ¹²⁵I-labeled fibrinogen was purchased from Amersham (UK). All other methods used in the present example have been previously described (22,27).

Construction of expression plasmids

The plasmids encoding SakSTAR(K35A,E38A,K74A,E75A), SakSTAR(E38A,E75A,R77A), SakSTAR(E38A,E75A), SakSTAR(K35A,E75A,R77A), SakSTAR(K35A,E75A), SakSTAR(E80A), SakSTAR(D82A), SakSTAR(E75A,D82A), SakSTAR(K74A) and SakSTAR(E75A) were constructed by the spliced overlap extension polymerase chain reaction (SOE-PCR) (24), using Vent DNA polymerase (New England Biolabs, Leusden, The Netherlands), and available or generated *sakSTAR* variants as template. Two fragments were amplified by PCR, the first one starting from the 5' end of the *staphylokinase* gene with primer 5'-CAGGAAACAGAATTCAGGAG-3' to the region to be mutagenized (forward primer), the second one from the same region (backward primer) to the 3' end of the *staphylokinase* gene with primer 5'-CAAAACAGCCAAGCTTCATTCATTCAGC-3'. The forward and backward primers shared an overlap of around 24 bp (primers not shown). The two purified fragments were then assembled together in a new primerless PCR using Taq polymerase (Boehringer Mannheim). After 7 cycles (1 min at 94°C, 1 min at 70°C), the extended product was reamplified by adding the 5' and 3' end primers (see above) to the PCR reaction and by cycling 25 times (1 min at 94°C, 1 min 55°C, 1 min at 72°C). The final product was purified, digested with EcoRI and HindIII and cloned into the corresponding sites of *pMEX602sakB*.

The plasmid encoding SakSTAR(E38A,K74A,E75A,R77A) was assembled by digestion of *pMEX602sakB* and *pMEX.SakSTAR(K35A,E38A,K74A,E75A,R77A)* with Bpm I which cuts between the codons for K35 and E38 of SakSTAR, and ligation of the required fragments. The plasmid encoding SakSTAR(K35A,K74A,E75A,R77A) was constructed by digestion of

pMEX.SakSTAR(K35A,E38A,K74A,E75A,R77A) and *pMEX.SakSTAR(K35A,E38A,E75A,R77A)* with Bpm I and religation of the required fragments. The plasmids encoding SakSTAR(K35A,E38A,E75A,R77A) and SakSTAR(K35A,E38A,K74A,R77A) were constructed by two PCR using *pMEX.SakSTAR(K35A,E38A,K74A,E75A,R77A)* as template, followed by restriction ligation and recloning into *pMEX602sakB*.

Expression and purification of SakSTAR variants

The SakSTAR variants were expressed and purified, as described below, from transformed *E. coli* WK6 grown either in LB medium [SakSTAR(E38A,K74A,E75A,R77A), SakSTAR(K74A), SakSTAR(E75A) and SakSTAR(E75A,D82A)], or in terrific broth (TB) (28) medium [SakSTAR(K35A,K74A,E75A,R77A), SakSTAR(K35A,E38A,E75A,R77A), SakSTAR(K35A,E38A,K74A,R77A), SakSTAR(K35A,E38A,E75A), SakSTAR(E38A,E75A,R77A), SakSTAR(E38A,E75A), SakSTAR(K35A,E75A,R77A), SakSTAR(K35A,E75A), SakSTAR(E80A), and SakSTAR(D82A)].

For derivatives produced in LB medium, a 20 mL aliquot of an overnight saturated culture was used to inoculate a 2 L volume of LB medium containing 100 µg/mL ampicillin. After 3 hours incubation at 37°C, IPTG (200 µmol/L) was added to induce expression from the *tac* promoter. The production phase was allowed to proceed for 4 hours, after which the cells were pelleted by centrifugation at 4,000 rpm for 20 min, resuspended in 1/20 volume (100 mL) of 0.01 mol/L phosphate buffer pH 6.5 and disrupted by sonication at 0°C. Cell debris were removed by centrifugation for 20 min at 20,000 rpm and the supernatant, containing the cytosolic soluble protein fraction, was stored at -20°C until purification.

For the derivatives produced in TB medium, a 4 mL aliquot of an overnight saturated culture in LB medium was used to inoculate a 2 L culture in terrific broth containing 100 µg/mL ampicillin. The culture was grown with vigorous aeration for 20 hours at 30°C. The cells were pelleted by centrifugation, resuspended in 1/10 volume (200 mL) of 0.01 mol/L phosphate buffer pH 6.5 and disrupted by sonication at 0°C. The suspension was then centrifuged for 20 min at 20,000 rpm and the supernatant was stored at -20°C until purification.

Cleared cell lysates containing the SakSTAR variants were subjected to chromatography on a 1.6 x 6 cm column of SP-Sephadex, followed by chromatography on a 1.6 x 5 cm column of Q-Sepharose [variants SakSTAR(E38A,K74A,E75A,R77A), SakSTAR(K35A,K74A,E75A,R77A), SakSTAR(K35A,E38A,E75A,R77A), SakSTAR(K35A,E38A,K74A,R77A) and SakSTAR(K35A,E38A,K74A,E75A)] or by chromatography on a 1.6 x 6 cm column of phenyl-Sepharose [variants SakSTAR(E35A,E38A,R77A), SakSTAR(E38A,E75A), SakSTAR(K35A,E75A,R77A), SakSTAR(K35A,E75A), SakSTAR(K74A), SakSTAR(E75A), SakSTAR(E80A), SakSTAR(D82A) and SakSTAR(E75A,D82A)]. The SakSTAR containing fractions, localized by SDS-gel electrophoresis, were pooled for further analysis.

Physicochemical and biochemical analysis

Protein concentrations were determined according to Bradford (29). The specific activities of SakSTAR solutions were determined with a chromogenic substrate assay carried out in microtiter plates using a mixture of 80 μ L SakSTAR solution and 100 μ L Glu-plasminogen solution prepared as described elsewhere (30) (final concentration 0.5 μ mol/L). After incubation for 30 min at 37°C, generated plasmin was quantitated by addition of 20 μ L S2403 (final concentration 1 mmol/L) and measurement of the absorption at 405 nm. The activity was expressed in home units (HU) by comparison with an in-house standard (lot STAN5) which was assigned an activity of 100,000 HU (100 kHU) per mg protein as determined by amino acid composition (7). SDS-PAGE was performed with the Phast SystemTM (Pharmacia, Uppsala, Sweden) using 10-15% gradient gels and Coomassie Brilliant blue staining. Reduction of the samples was performed by heating at 100°C for 3 min in the presence of 1% SDS and 1% dithioerythritol. The specific activities of the different SakSTAR mutants determined with the chromogenic substrate assay are summarized in Table 1.

Binding to murine monoclonal antibodies

In agreement with previous observations (22), SakSTAR(K74A,E75A,R77A) did not react with 4 of the 5 MAbs recognizing epitope I, whereas SakSTAR(K35A,E38A) did not react with 3 of the 5 and SakSTAR(E80A,D82A) not with 4 of the 5 MAbs recognizing epitope III. These reduced reactivities were additive in SakSTAR(K35A,E38A,K74A,E75A,R77A) and in SakSTAR(K74A,E75A,R77A,E80A,D82A). The reduced reactivity of SakSTAR(K74A,E75A,R77A) was fully maintained in SakSTAR(K35A,E38A,K74A,E75A) and in SakSTAR(K35A,E75A,R77A), largely in SakSTAR(K35A,E38A,E75A,R77A), SakSTAR(E38A,E75A,R77A), SakSTAR(E38A,E75A) and SakSTAR(E75A), but much less in SakSTAR(K35A,E38A,K74A,R77A) and SakSTAR(K74A), indicating that E75 is the main contributor to the binding of the 4 MAbs recognizing epitope I of SakSTAR. However, surprisingly, binding of epitope I antibodies to SakSTAR(E75A,D82A) was normal in two independent preparations from expression plasmids with confirmed DNA sequences. The reduced reactivity of the 3 MAbs of epitope III with SakSTAR(K35A,E38A) required both K35 and E38, as demonstrated with SakSTAR(E38A,K74A,E75A,R77A) and SakSTAR(K35A,K74A,E75A,R77A), with SakSTAR(E38A,E75A) and SakSTAR(K35A,E75A) and with SakSTAR(E38A,E75A,R77A) and SakSTAR(K35A,E75A,R77A). The reduced reactivity of the 4 MAbs of cluster III with SakSTAR(E80A,D82A) was maintained in SakSTAR(D82A) but not in SakSTAR(E80A).

Absorption of antibodies, elicited in patients by treatment with wild-type SakSTAR

Plasma samples from 16 patients with acute myocardial infarction, obtained several weeks after treatment with SakSTAR (4,31) were used. The staphylokinase-neutralizing activity in these samples was determined as follows. Increasing concentrations of wild-type or variant SakSTAR (50 μ L volumes containing 0.2 to 1000 μ g/mL) were added to a mixture of 300 μ L citrated human plasma and 50 μ L buffer or test plasma, immediately followed by addition of 100 μ L of a mixture containing thrombin (50 NIH units/mL) and CaCl_2 (25 mmol/L). The plasma clot lysis time was measured and plotted against the concentration of SakSTAR moiety. From this curve the concentration of staphylokinase moiety that produced complete

clot lysis in 20 min was determined. The neutralizing activity titer determined as the difference between the test plasma and buffer values and was expressed in μg per mL test plasma. The results of the individual patients have been reported elsewhere (22). For the present invention, three plasma pools were made, one from 10 patients from whom sufficient residual plasma was available, one from three patients that absorbed less than 50% of the antibodies with SakSTAR(K35A,E38A,K74A,E75A,R77A) (Subpool B) and one from three patients that absorbed >90% of the antibodies with SakSTAR(K35A,E38A,K74A,E75A,R77A) (Subpool C).

These plasma pools were diluted (1/30 to 1/200) until their binding to SakSTAR substituted chips in the BIAcore instrument amounted to approximately 2000 RU. From this dilution a calibration curve for antibody binding was constructed using further serial two-fold dilutions. The plasma pools were absorbed for 10 min with 100 nmol/L of the SakSTAR variants, and residual binding to immobilized SakSTAR was determined. Residual binding was expressed in percent of unabsorbed plasma, using the calibration curve.

The results are summarized in Table 1. Whereas wild-type SakSTAR absorbed more than 95% of the binding antibodies from pooled plasma of the 10 patients, incomplete absorption (<60%) was observed with SakSTAR(K74A,E75A,R77A), SakSTAR(K35A,E38A,K74A,E75A,R77A), SakSTAR(E38A,K74A,E75A,R77A), SakSTAR(K35A,K74A,E75A,R77A), SakSTAR(K35A,E38A,K74A,R77A), SakSTAR(K35A,E38A,K74A,E75A), SakSTAR(K74A) and SakSTAR(K74A,E75A,R77A,E80A,D82A) but absorption was nearly complete with SakSTAR(K35A,E38A), SakSTAR(K35A,E38A,E75A,R77A), SakSTAR(E38A,E75A,R77A), SakSTAR(E38A,E75A), SakSTAR(K35A,E75A,R77A), SakSTAR(K35A,E75A), SakSTAR(E75A), SakSTAR(E80A,D82A), SakSTAR(E80A), SakSTAR(D82A) and SakSTAR(E75A,D82A). These results, surprisingly, demonstrate that approximately 40% of the antibodies elicited in patients by treatment with wild-type SakSTAR depend on K74 for their binding (Table 1). Absorption with pooled plasma from 3 patients from which <50% of the antibodies were absorbed with SakSTAR(K35A,E38A,K74A,E75A,R77A) (Subpool B) confirmed the predominant role of K74 for antibody recognition. As expected, absorption with pooled plasma from 3 patients

from which >95 of the antibodies were adsorbed with SakSTAR(K35A,E38A,K74A,E75A,R77A) (Subpool C) was nearly complete with all variants tested.

EXAMPLE 3

Comparative thrombolytic efficacy and immunogenicity of SakSTAR(K74A,E75A,R77A) and SakSTAR(K74A) versus SakSTAR in patients with peripheral arterial occlusion

Purification of SakSTAR(K74A,E75A,R77A) and SakSTAR(K74A) for use in vivo

A 12 to 24 L culture (in 2 L batches) of the variants SakSTAR(K74A,E75A,R77A), or of SakSTAR(K74A) was grown and IPTG-induced in LB medium supplemented with 100 µg/mL ampicillin, pelleted, resuspended, disrupted by sonication and cleared as described above. The compounds were purified by chromatography on a 5 x 20 cm column of SP-Sephadex, a 5 x 10 cm column of Q-Sepharose and/or a 5 x 13 cm column of phenyl-Sepharose using buffer systems described elsewhere (22,23). The materials were then gel filtered on sterilized Superdex 75 to further reduce their endotoxin content. The SakSTAR variant containing fractions were pooled, the protein concentration was adjusted to 1 mg/mL and the material sterilized by filtration through a 0.22 µm Millipore filter. The methodology used to determine the biological properties of the final material required for use in vivo is described above and elsewhere (22).

Materials and Methods

Staphylokinase-neutralizing activity in plasma was determined as described above. Quantitation of antigen-specific IgG and IgM antibodies was performed using enzyme-linked immunosorbent assays in polystyrene microtiter plates essentially as described previously (22). In the IgG assays, dilution curves of affinospecific anti-SakSTAR IgG antibodies were included on each plate. These antibodies were isolated from plasma obtained from 3 patients, after thrombolytic therapy with wild-type SakSTAR, by chromatography on protein A-

Sepharose and on insolubilized SakSTAR, and elution of bound antibodies with 0.1 mol/L glycine-HCl, pH 2.8. The purity of the IgG preparation was confirmed by sodium dodecylsulfate polyacrylamide gel electrophoresis. In the IgM assays, titers defined as the plasma dilution giving an absorbancy at 492 nm equivalent to that of a 1/640 dilution of pooled plasma were determined and compared with the titer of baseline samples before treatment (median value 1/410, interquartile range 1/120-1/700).

Thrombolytic efficacy

Wild-type SakSTAR or the variants SakSTAR(K74A) or SakSTAR(K74A,E75A,R77A) were administered intra-arterially at or in the proximal end of the occlusive thrombus as a bolus of 2 mg followed by an infusion of 1 mg/hr (reduced overnight in some patients to 0.5 mg/hr) in groups of 6 to 12 patients with angiographically documented occlusion of a peripheral artery or bypass graft of less than 120 days duration. Patients were studied after giving informed consent, and the protocol was approved by the Human Studies Committee of the University of Leuven. Inclusion and exclusion criteria, conjunctive antithrombotic treatment (including continuous intravenous heparin) and the study protocol were essentially as previously described (22).

Relevant baseline characteristics of the individual patients are shown in Table 2. The majority of PAO were at the femoropopliteal level. Two iliac stent and 8 graft occlusions were included. Eight patients presented with incapacitating claudication, 5 with chronic ischemic rest pain, 7 with subacute ischemia and 7 with acute ischemia. One patient (POE) who had 2 years previously been treated with SakSTAR was included in the SakSTAR(K74A) group. This patient was not included in the statistical analyses.

Table 2 also summarizes the individual treatment and outcome. Intra-arterial infusion, at a dose of 6.0 to 25 mg and a duration of 4.0 to 23 hrs, induced complete recanalization in 24 patients and partial recanalization in 3. Complementary endovascular procedures (mainly PTA) were performed in 17 patients and complementary reconstructive vascular surgery following thrombolysis in 3. No patient underwent major amputation. Early recurrence of thrombosis after the end of the angiographic procedure occurred in 4 patients. Bleeding complications were absent or limited to mild to moderate hematoma formation at the

angiographic puncture site except for 5 patients who required transfusion (data not shown). Intracranial or visceral hemorrhage was not observed. Circulating fibrinogen, plasminogen and α_2 -antiplasmin levels remained essentially unchanged during infusion of the SakSTAR moieties (data not shown), confirming absolute fibrin specificity of staphylokinase at the dosages used. Significant in vivo fibrin digestion occurred as evidenced by elevation of fibrin fragment D-dimer levels. Intra-arterial heparin therapy prolonged aPTT levels to a variable extent (data not shown).

Antibody induction

Antibody-related SakSTAR-, SakSTAR(K74A)- and SakSTAR(K74A,E75A,R77A)-neutralizing activity and anti-SakSTAR, anti-SakSTAR(K74A) and anti-SakSTAR(K74A,E75A,R77A) IgG, were low at baseline and during the first week after the infusion (Figure 2). From the second week on, neutralizing activity levels increased to reach median values at 3 to 4 weeks of 20 μ g SakSTAR(K74A) and 2.4 μ g SakSTAR(K74A,E75A,R77A) neutralized per mL plasma in the patients treated with SakSTAR(K74A) and SakSTAR(K74A,E75A,R77A), respectively, which is significantly lower than the median value of 93 μ g wild-type SakSTAR neutralized per mL in the patients treated with SakSTAR ($p = 0.024$ for differences between the three groups by Kruskal-Wallis analysis and $p = 0.01$ and $p = 0.036$, respectively, for variants vs wild-type by Mann-Whitney rank sum test). The levels of anti-SakSTAR(K74A) and of anti-SakSTAR(K74A,E75A,R77A) IgG increased to median values at 3 to 4 weeks of 270 and 82 μ g/mL plasma in patients treated with SakSTAR(K74A) and SakSTAR(K74A,E75A,R77A) respectively, which is significantly lower than the median value of 1800 μ g anti-SakSTAR per mL plasma in the patients treated with SakSTAR ($p = 0.024$ for differences between the three groups by Kruskal-Wallis analysis and $p = 0.007$ and 0.05 , respectively, for variants versus wild-type by Mann-Whitney rank sum test).

The titers of anti-SakSTAR(K74A) and of anti-SakSTAR(K74A,E75A,R77A) IgM increased from median baseline values of 1/460 and 1/410 to median values at 1 week of 1/510 and 1/450 in patients treated with SakSTAR(K74A) and SakSTAR(K74A,E75A,R77A), respectively, which was not significantly different from the median values of 1/320 at

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baseline and 1/640 at week 2 in patients treated with SakSTAR. Corresponding values at 2 weeks were 1/590 and 1/550 in patients given SakSTAR(K74A) and SakSTAR(K74A,E75A,R77A), not significantly different from 1/930 with SakSTAR (data not shown).

The antibodies induced by treatment with SakSTAR were completely absorbed by SakSTAR but incompletely by SakSTAR(K74A) and by SakSTAR(K74A,E75A,R77A) confirming the immunogenicity of the K74,E75,R77 epitope and the dominant role of K74 in the binding of antibodies directed against this epitope. The antibodies induced by treatment with SakSTAR(K74A) or SakSTAR(K74A,E75A,R77A) were completely absorbed by SakSTAR, by SakSTAR(K74A) and by SakSTAR(K74A,E75A,R77A), indicating that immunization was not due to neoepitopes generated by substitution of Lys74 with Ala, but to epitopes different from the K74,E75,R77 epitope.

Thus, this example illustrates that staphylokinase variants with *reduced* antibody induction but *intact* thrombolytic potency can be generated. The present experience in 26 patients treated with SakSTAR (n= 9), SakSTAR(K74A) (n= 11) and SakSTAR(K74A,E75A,R77A) (n= 6) combined with previous experience in 14 patients with SakSTAR (n= 7) and SakSTAR(K35A,E38A,K74A,E75A,R77A) (n= 7) (31) and in 24 patients with SakSTAR (32), and with subsequent non-randomized experience in patients with SakSTAR (n= 30) with SakSTAR(K74A) (n= 12) and with SakSTAR(K74A,E75A,R77A) (n= 7) (data not shown), allows an initial estimation of the prevalence of immunization by intra-arterial treatment with SakSTAR or variants with an altered K74,E75,R77 epitope [SakSTAR(K74A), SakSTAR(K74A,E75A,R77A) and SakSTAR(K35A,E38A,K74A,E75A,R77A)]. Neutralizing activity data after 2 to 4 weeks, available in 70 patients with peripheral arterial occlusion given intra-arterial SakSTAR, revealed that 56 patients (80 percent) had levels > 5 µg compound neutralized per mL plasma. Of the patients given SakSTAR(K74A), SakSTAR(K74A,E75A,R77A) or SakSTAR(K74A,E75A,K74A,E75A,R77A), 27 of the 43 (63 percent) had neutralizing activity levels of > 5 µg compound per mL plasma. This difference is statistically significant (p= 0.05 by Fisher's exact test) indicating that the K74,E75,R77 epitope is a major determinant of antibody induction. However, the residual

prevalence of specific immunocompetence against SakSTAR(K74A) indicates that additional mutagenesis to further reduce the immunogenicity of SakSTAR variants for clinical use, would be desirable.

EXAMPLE 4

Construction, epitope mapping with murine monoclonal antibodies and absorption with pooled plasma of immunized patients, of alanine-substitution mutants of staphylokinase

Site-directed mutagenesis was applied to residues other than "charged amino acids" in order to identify i) additional residues belonging to epitopes I and III identified with the panel of murine Mabs and ii) amino acids determining absorption to antiserum from immunized patients. Since functional epitopes generally comprise more than one amino acid residue critical for antibody binding, identification of additional residues in these epitopes could lead to the construction of new combination derivatives displaying a lower antigenic profile, while keeping the specific activity and the temperature stability of wild-type staphylokinase.

In this example, the construction and characterization of SakSTAR variants in which one or at most two amino acids (adjacent or in close vicinity) were substituted with alanine is described. The mutants described under this example are listed in Table 3. These variants were expressed in *E. coli*, purified and characterized in terms of specific activity, reactivity with the panel of murine monoclonal antibodies, and absorption of antibodies from plasma of patients treated with wild-type SakSTAR.

Reagents and Methods

The source of all reagents used in the present study has previously been reported (22), or is specified below. The template vector for mutagenesis, *pMEX602sakB* (i.e. *pMEX.SakSTAR*), has been described elsewhere (23). Restriction and modification enzymes were purchased from New England Biolabs (Leusden, The Netherlands), Boehringer Mannheim (Mannheim, Germany) or Pharmacia (Uppsala, Sweden). The enzymatic reactions were performed according to the supplier recommendation. The mutagenic oligonucleotides and primers were obtained from Eurogentec (Seraing, Belgium). Plasmid DNA was isolated using a

Construction of expression plasmids

The variants SakSTAR(I128A), SakSTAR(L127A) and SakSTAR(N126V) were constructed by performing a polymerase chain reaction using the primer 818A located at the 5' end of the *sakSTAR* gene and mutagenic primers (not shown). The amplified product (30 cycles: 1 sec at

94°C, 1 sec at 50°C, 10 sec at 72°C) was purified, digested with EcoRI and StyI, and ligated into the corresponding sites of *pMEXSakSTAR*. The variant SakSTAR(F125A) was constructed by performing two consecutive PCR reactions (30 cycles: : 30 sec at 94°C, 30 sec at 50°C, 30 sec at 72°C). In the first reaction, a fragment of *pMEX.SakSTAR* was amplified with the primers 818A and a mutagenic primer. This amplified fragment was then used as template in a second PCR reaction with a mutagenic primer in order to further elongate the fragment downstream of the *StyI* site present in the *sakSTAR* gene (corresponding to amino acids 130-131 of SakSTAR). The resulting product was digested with EcoRI and StyI, and ligated into the corresponding sites of *pMEXSakSTAR*.

The plasmids encoding all the other variants listed in Table 3 were constructed by direct PCR or by the spliced overlap extension polymerase chain reaction (SOE-PCR)(24) using *pMEX.SakSTAR* or available plasmids encoding SakSTAR variants as template. Two fragments were amplified by PCR (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C), the first one starting from the 5' end (primer 818A) of the *staphylokinase* gene to the region to be mutagenized (forward primer), the second one from this same region (backward primer) to the 3' end of the gene with primer 818D (5' CAAACAGCCAAGCTTCATTCATTCAGC). The forward and backward primers shared an overlap of around 24 bp (primers not shown). The two purified fragments were then assembled together in a second PCR reaction with the external primers 818A and 818D (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C). The amplified product from this final reaction was purified, digested with EcoRI and HindIII and ligated into the corresponding site of *pMEX.SakSTAR*. For each construction, the sequence of the variant was confirmed by sequencing the entire *SakSTAR* coding region.

Expression and purification of SakSTAR variants

The SakSTAR variants were expressed and purified, as described below, from transformed *E. coli* grown in terrific broth (TB) medium (28). A 2 to 4 mL aliquot of an overnight saturated culture in LB medium was used to inoculate a 1 to 2 L culture in terrific broth supplemented with 100 µg/mL ampicillin. The culture was incubated with vigorous aeration

and at 30°C. After about 16 hours incubation, IPTG (200 μ mol/L) was added to the culture to induce expression from the *tac* promoter. After 3 hours induction, the cells were pelleted by centrifugation at 4,000 rpm for 20 min, resuspended in 1/10 volume of 0.01 mol/L phosphate buffer pH 6-6.5 and disrupted by sonication at 0°C. The suspension was centrifuged for 20 min at 20,000 rpm and the supernatant was stored at 4°C or at -20°C until purification. The material was purified essentially as described above (Example 2): cleared cell lysates containing the SakSTAR variants were subjected to chromatography on a 1.6 x 5 cm column of SP-Sephadex, followed by chromatography on a 1.6 x 8 cm column of phenyl-Sepharose. The SakSTAR containing fractions, localized by SDS-gel electrophoresis, were pooled for further analysis.

Physicochemical and biochemical analysis

Protein concentrations were determined according to Bradford (29). SDS-PAGE was performed with the Phast System™ (Pharmacia, Uppsala, Sweden) using 10-15% gradient gels and Coomassie Brilliant blue staining, and the specific activities of SakSTAR solutions were determined with a chromogenic substrate assay carried out in microtiter plates (as described in example 2). The specific activity of the different SakSTAR variants are summarized in Table 3.

Reactivity of SakSTAR variants with a panel of murine monoclonal antibodies

The methodology used to determine the reactivity of the SakSTAR variants with a panel of murine monoclonal antibodies was described in example 1 above. The results are summarized in Table 3 (the layout of this Table corresponds to the layout of Table 1, as described in example 1). Apparent association constants at least 10-fold lower than those of wild-type staphylokinase were considered as significant and are indicated in bold type in the table.

In order to obtain a comprehensive inventory of the properties of Ala-substitution variants of the SakSTAR molecule, 67 plasmids encoding variants with substitution of a single or two adjacent amino acids with Ala were constructed, expressed and purified. Together with the 35

charged residue to Ala-substitution variants previously described (22, and example 2), this analysis covers all residues in SakSTAR except Gly, Ala and Pro, as illustrated in Figure 3. Eight of the variants could not be obtained in purified form, primarily as a result of low expression levels, 11 variants were inactive, 56 had a reduced specific activity, and 27 had a maintained or increased specific activity (≥ 100 kHU/mg). The yields of purified material from cultures of expressed plasmids were 16 mg/L (median, 10 to 90 percentile range 4 to 41 mg/L). SDS polyacrylamide gel electrophoresis consistently showed one main band with $M_r \approx 16,000$, usually representing $\geq 95\%$ of total protein (not shown).

Substitution of K35, N95, S103 or K135 with Ala yielded variants with specific activities of ≥ 200 kU/mg. Substitution of W66, Y73 or E75 with Ala reduced the reactivity of the variants with ≥ 3 antibodies of epitope cluster I, of H43 or V45 with Ala that with 3 antibodies from epitope cluster II and of V32, K35, D82 and K130 with Ala that with ≥ 3 antibodies of epitope cluster III.

Absorption of antibodies, elicited in patients by treatment with SakSTAR

For the present example, the three plasma pools, as described in example 2 were used. The methodology used to evaluate the absorption with wild-type staphylokinase and with SakSTAR variants, of antibodies elicited in patients treated with SakSTAR, is described in detail in example 2. The results are summarized in Table 3. Whereas wild-type SakSTAR and most of the variants analyzed in this example absorbed more than 95% of the binding antibodies from pooled plasma of the 10 patients, incomplete absorption ($<60\%$) was observed with SakSTAR(Y73A), and with SakSTAR(K74A). The predominant role of Lys74 for antibody recognition has been demonstrated previously (see example 2). The present results indicate that Tyr73 participates to the same major epitope as Lys74, or, alternatively, that substitution at Tyr73 may indirectly induce a structural modification of the "K74-epitope". Absorption with pooled plasma from 3 patients from which $>95\%$ of the antibodies were absorbed with SakSTAR(K35A,E38A,K74A,E75A,R77A) (Subpool C, see example 2) was nearly complete with most variants tested.

EXAMPLE 5

Construction, epitope mapping with murine monoclonal antibodies and absorption with pooled plasma of immunized patients, of staphylokinase variants with substitution of S34, G36 and/or H43

The natural variant Sak42D differs from SakSTAR in three amino acids and corresponds to SakSTAR(S34G,G36R,H43R). Sak42D is characterized by reduced reactivity with some murine antibodies of epitope clusters II and III and a slightly reduced absorption of antibodies from plasma of patients treated with SakSTAR (Table 4). Mutagenesis of these residues in SakSTAR revealed that the reduced reactivity with epitope cluster III and with immunized patient plasma could be ascribed to the G36R substitution, the H43R substitution mediated the reduced reactivity with epitope cluster II but had no effect on the reactivity with immunized patient plasma, whereas the S34A substitution had no effect. The G36R substitution could be combined with the K74R but not with the K74A substitution, without significant reduction of the specific activity (Table 4).

EXAMPLE 6

Construction, epitope mapping with murine monoclonal antibodies and absorption with pooled plasma of immunized patients, of staphylokinase variants with substitution of K35, E65, Y73, K74, E80+D82, N95, K130, V132 and/or K135

Based on the results of the alanine-substitution analysis in example 4, K35, N95 and K135 were selected for further analysis because SakSTAR(K35A), SakSTAR(N95A) and SakSTAR(K135A) had a two-fold increased specific activity, Y73 and K74 because SakSTAR(Y73A) and SakSTAR(K74A) had a markedly reduced reactivity with antibodies from epitope cluster I and diminished absorption of antibodies from plasma of patients immunized by treatment with SakSTAR, and K35, E80+D82, K130 and V132 because SakSTAR(K35A), SakSTAR(E80A,D82A), SakSTAR(K130A) and SakSTAR(V132A) had a reduced reactivity with antibodies from epitope cluster III.

In an effort to maximize the activity/antigenicity ratio, these amino acids were substituted with other amino acids than Ala. As summarized in Table 5, substitution of K35 with A, E or Q revealed that SakSTAR(K35A) had the most interesting properties, substitution of Y73 with F, H, L, S or W did not rescue the marked reduction in specific activity, and K74 confirmed its key role in binding of antibodies from immunized patient plasma, the best specific activity/antigenicity ratios being obtained with SakSTAR(K74Q) and SakSTAR(K74R). SakSTAR(E80A,D82A) was preferred over the single residue variants SakSTAR(E80A) or SakSTAR(D82A) because of its somewhat lower reactivity with immunized patient plasma. SakSTAR(N95A) could not be further improved by substitution of N95 with E, G, K or R and it was unable to confer its increased specific activity to variants containing K74A or K135R. Finally SakSTAR(K130A) was outperformed in terms of specific activity by SakSTAR(K130T) and SakSTAR(V132A) by SakSTAR(V132R).

EXAMPLE 7

Construction, epitope mapping with murine monoclonal antibodies and absorption with pooled plasma of immunized patients of combination variants of SakSTAR(K130T,K135R) and SakSTAR(E80A,D82A,K130T,K135R) with K35A,G36R,E65X,K74X and selected other amino acids

In the present and the following examples an additional plasma pool was made from 40 patients obtained several weeks after treatment with SakSTAR (Pool 40). The original pool from 10 patients is further identified as Pool 10. The absorption of staphylokinase-specific antibodies was quantified as described above and elsewhere (22).

The SakSTAR(K130T,K135R) variant was taken as a template for additive mutagenesis because of its high specific activity with a moderate reduction of binding to antibodies of epitope cluster III and absorption of antibodies from immunized patient plasma (Table 6). Addition of G36R, K74R, or K74Q or both to the template did not markedly reduce the specific activity, reduced the reactivity with monoclonal antibodies against epitope cluster III (G36R substitution) and decreased the absorption of antibodies from immunized patient plasma (K74R or K74Q substitution). Combination of E65A or E65Q with K74Q in the

Combination of the SakSTAR(E80A,D82A) and SakSTAR(K130T,K135R) templates, did not affect the specific activity and had a reduced reactivity with epitope cluster III antibodies (Table 7). Therefore the SakSTAR(E80A,D82A,K130T,K135R) template was selected for further mutagenesis. Addition of K74R and even more of K74Q drastically reduced the reactivity with immunized patient plasma. Finally, addition of E65D or of K35A or E65S to the SakSTAR(K74R,E80A,D82A,K130T,K135R) or SakSTAR(K74Q,E80A,D82A,K130T,K135R) templates yielded variants with intact specific activity which only bound ≤ 45 of the antibodies of pooled immunized patient plasma and less than 15 percent of the subpool reacting for more than 50 percent with the K74,E75,R77 epitope.

Characterization of selected variants of staphylokinase with intact specific activity and less than 50% adsorption of pooled SakSTAR specific human antibodies elicited in patients by treatment with wild-type SakSTAR.

Twenty three of the variants constructed and characterized in the above examples combined the properties of a residual specific activity of ≥ 100 kHU/mg and ≤ 50 percent absorption with the pool of antisera obtained from 10 patients treated with wild-type SakSTAR. The results are summarized in Table 8. Results obtained with Subpool B and Subpool C and with the pool of 40 patients treated with wild-type SakSTAR are included. SakSTAR(K74Q,E80A,D82A,K130T,K135R), SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R), SakSTAR(K35A,E65D,K74Q,E80A,D82A,K130T,K135R) and SakSTAR(E65Q,

K74Q,N95A,E118A,K130A,K135R,K136A,+137K) were selected for further characterization.

Fibrinolytic properties of SakSTAR variants in human plasma in vitro

The fibrinolytic and fibrinogenolytic properties of the SakSTAR variants were determined as previously described. Dose- and time-dependent lysis of ^{125}I -fibrin labeled human plasma clots submerged in human plasma was obtained with the selected variants (Table 9). Spontaneous clot lysis during the experimental period was $\leq 5\%$ (not shown). Equi-effective concentrations of test compound (causing 50% clot lysis in 2 hrs; C_{50}), determined graphically from plots of clot lysis at 2 hrs versus the concentration of plasminogen activator (not shown), ranged from 0.11 ± 0.01 to 0.24 ± 0.04 $\mu\text{g/mL}$ at which the residual fibrinogen levels ranges between 92 ± 30 and 97 ± 30 percent of baseline (Table 9). The concentrations of compound causing 50% fibrinogen degradation in 2 hrs in human plasma in the absence of fibrin were determined graphically from dose-response curves (not shown). These values (mean \pm SD of 3 independent experiments) ranged from 14 ± 3.2 to 29 ± 3.1 $\mu\text{g/mL}$ (Table 9). Surprisingly the very high specific activity of SakSTAR(E65Q,K74Q,N95A,E118A,K130A,K135R,K136A,+137K) in the chromogenic assay was not associated with an increased thrombolytic potency in a plasma milieu.

Temperature stability of selected SakSTAR variants

The temperature stability of preparations of SakSTAR(K74Q,E80A,D82A,K130T,K135R), SakSTAR(E65D,K74R,E80A, D82A,K130T,K135R) and SakSTAR(K35A,E65D,K74Q,E80A,D82A,K130T,K135R), dissolved to a concentration of 1.0 mg/mL in 0.15 mol/L NaCl, 0.01 mol/L phosphate buffer, pH 7.5 at various temperatures is illustrated in Fig. 4. At temperatures up to 37°C , all compounds remained fully active for up at least three days. At 56°C and 70°C the three variants were however less stable than wild-type SakSTAR.

Pharmacokinetic properties of SakSTAR variants following bolus injection in hamsters

The pharmacokinetic parameters of the disposition of SakSTAR variants from blood were evaluated in groups of 4 hamsters following intravenous bolus injection of 100 µg/kg SakSTAR variant. SakSTAR-related antigen was assayed using the ELISA described elsewhere. The ELISA was calibrated against each of the SakSTAR variants to be quantitated. Pharmacokinetic parameters included: initial half-life (in min), $t_{1/2\alpha} = \ln 2/\alpha$; terminal half-life (in min), $t_{1/2\beta} = \ln 2/\beta$; volume of the central (plasma) compartment (in mL), $V_C = \text{dose}/(A+B)$; area under the curve (in µg.min.mL⁻¹), $AUC = A/\alpha + B/\beta$; and plasma clearance (in mL.min⁻¹), $Cl_p = \text{dose}/AUC$ (33).

The disposition rate of staphylokinase-related antigen from blood following bolus injection of 100 µg/kg of the selected SakSTAR variants in groups of 4 hamsters could adequately be described by a sum of two exponential terms by graphical curve peeling (results not shown), from which the pharmacokinetic parameters summarized in Table 10 were derived. The pharmacokinetic parameters of the mutants were not markedly different from those of wild type SakSTAR. Initial plasma half-lives ($t_{1/2}(\alpha)$) ranged between 2.0 and 3.2 min and plasma clearances (Cl_p) between 1.6 and 4.1 mL/min.

EXAMPLE 9

Comparative thrombolytic efficacy and immunogenicity of SakSTAR(K74Q, E80A, D82A, K130T, K135R) and SakSTAR(E65D, K74R, E80A, D82A, K130T, K135R) versus SakSTAR in patients with peripheral arterial occlusion

Purification for use in vivo

Eighteen liter cultures (in 2 L batches) of the variants SakSTAR(K74Q, E80A, D82A, K130T, K135R) and SakSTAR(E65D, K74R, E80A, D82A, K130T, K135R) were grown for 20 hours in terrific broth medium (28), supplemented with 100 µg/mL ampicillin and induced with IPTG during the last 3 hours. The cells were pelleted, resuspended in 1/10 volume of 0.01 mol/L phosphate buffer, pH 6.0, disrupted by sonication and cleared by centrifugation. The

Out of culture volumes of 18 liters of SakSTAR variant, 840 mg of SakSTAR(K74Q,E80A,D82A,K130T,K135R) with a specific activity of 140 kHU/mg and 800 mg SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) with a specific activity of 150 were purified. The endotoxin content was <0.1 and 0.26 IU/mg. Gel filtration on HPLC revealed a single main symmetrical peak in the chromatographic range of the column, representing >98% of the eluted material (total area under the curve) (not shown). SDS gel electrophoresis of 40 µg samples revealed single main components (not shown). Preparations sterilized by filtration proved to be sterile on 3 day testing as described elsewhere (22). Intravenous bolus injection of SakSTAR variants in groups of 5 mice (3 mg/kg body weight), did not provoke any acute reaction, nor reduced weight gain within 8 days, in comparison with mice given an equal amount of saline (not shown).

Wild-type SakSTAR or the variants SakSTAR(K74Q,E80A,D82A,K130T,K135R) or SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) were administered intra-arterially at or

Relevant baseline characteristics of the individual patients and results of treatment and outcome are shown in Table 11. Intra-arterial infusion, at a dose of 3.5 to 27 mg and a duration of 2 to 44 hrs, induced complete recanalization in 22 patients and partial recanalization in 5. Complementary endovascular procedures (mainly PTA) were performed in 13 patients and complementary reconstructive vascular surgery following thrombolysis in 5. One patient underwent major amputation. Bleeding complications were usually absent or limited to mild to moderate hematoma formation at the angiographic puncture sites (data not shown). One patient, given wild-type SakSTAR suffered a non-fatal intracranial bleeding, one (BUE) a retroperitoneal hematoma and two (MAN and STRO) a gastro-intestinal bleeding.

Circulating fibrinogen, plasminogen and α_2 -antiplasmin levels remained unchanged during infusion of the SakSTAR moieties (data not shown), reflecting absolute fibrin specificity of these agents at the dosages used (data not shown). Significant in vivo fibrin digestion occurred as evidenced by elevation of fibrin fragment D-dimer levels. Intra-arterial heparin therapy prolonged aPTT levels to a variable extent (data not shown).

Staphylokinase-neutralizing activity in plasma and antigen-specific IgG antibodies were quantitated essentially as described above and elsewhere (22).

Antibody-related SakSTAR-, SakSTAR(K74Q,E80A,D82A,K130T,K135R)- and SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)-neutralizing activity and anti-SakSTAR, anti-SakSTAR(K74Q,E80A,D82A,K130T,K135R) and anti-

SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) IgG, were low at baseline and during the first week after the infusion (Figure 5). From the second week on, neutralizing activity levels increased to reach median values at 3 to 4 weeks of 9 µg SakSTAR(K74Q,E80A,D82A,K130T,K135R) and 0.5 µg SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) neutralized per mL plasma in the patients treated with the corresponding moieties, respectively, as compared to median value of 24 µg wild-type SakSTAR neutralized per mL in the 15 patients treated with SakSTAR. The levels of anti-SakSTAR(K74Q,E80A,D82A,K130T,K135R) and of anti-SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) IgG increased to median values at 3 to 4 weeks of 420 and 30 µg/mL plasma in patients treated with the corresponding moieties, respectively, as compared to a median value of 590 µg anti-SakSTAR per mL plasma in the patients treated with SakSTAR (Figure 5). The prevalence of immunization, defined as neutralizing activities in plasma after 2 to 4 weeks exceeding 5 µg/ml was 3 of 6 patients (50 percent) with SakSTAR(K74Q,E80A,D82A,K130T,K135R), 1 of 6 patients (17 percent) with SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R), as compared to 56 of 70 patients (80 percent) with SakSTAR. This difference is statistically highly significant ($p=0.01$ by 2×3 Chi square analysis).

The antibodies induced by treatment with SakSTAR were completely absorbed by SakSTAR but incompletely by SakSTAR(K74Q,E80A,D82A,K130T,K135R) and by SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) (Table 12). Antibodies induced by treatment with SakSTAR(K74Q,E80A,D82A,K130T,K135R), detectable in 4 of the 6 patients, were completely (≥ 90 percent) absorbed by SakSTAR, by SakSTAR(K74Q,E80A,D82A,K130T,K135R) and by SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R), indicating that immunization was not due to neopeptides generated by substitution of wild-type amino acids. Antibodies induced by treatment with SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) detectable in one patient (URB) were completely absorbed with SakSTAR(K74Q,E80A,D82A,K130T,K135R) and with SakSTAR(E65D,K74Q,E80A,D82A,K130T,K135R) but incompletely (85%) with wild-type SakSTAR, suggesting that a small fraction of the induced antibodies might be directed against a neopeptide in the variant used for infusion.

EXAMPLE 10

Construction, purification and characterization of cysteine-substitution mutants of staphylokinase

Site-directed mutagenesis was applied to substitute exposed amino acids with single cysteine residues in order to construct i) homodimeric forms of staphylokinase, upon formation of an intermolecular disulfide bridge, and ii) polyethylene glycol-conjugated molecules (PEG-derivatives). The aim for this study is twofold: first, the clearance can be reduced by increasing the size of the injected molecule (via dimerization or conjugation with large molecule such as PEG) and second, PEG-derivatives have also been shown to induce a reduced immunoreactivity in animal models (for review, see ref. 34). In both cases, a prolonged half-life *in vivo* could help to reduce the pharmacological dosis of straphylokinase in patients. This reduction could be accompanied with a reduced immunogenic reaction against the thrombolytic agent.

In this example, the construction and characterization of two SakSTAR variants in which one single amino acid was substituted with cysteine is described. The mutants described under this example are listed in Table 13. These variants were expressed in *E. coli*, purified and characterized to some extent in terms of specific activity, fibrinolytic properties in human plasma *in vitro* and pharmacokinetic properties following bolus injection in hamsters.

Reagents and Methods

The source of all reagents used in the present study has previously been reported (22), or is specified below. The template vector for mutagenesis, *pMEX602sakB* (i.e. *pMEX.SakSTAR*), has been described elsewhere (23). Restriction and modification enzymes were purchased from New England Biolabs (Leusden, The Netherlands), Boehringer Mannheim (Mannheim, Germany) or Pharmacia (Uppsala, Sweden). The enzymatic reactions were performed according to the supplier recommendation. The mutagenic oligonucleotides and primers were obtained from Eurogentec (Seraing, Belgium). Plasmid DNA was isolated using a purification kit from Qiagen (Hilden, Germany), as recommended. Transformation-competent *E. coli* cells were prepared by the well-known calcium phosphate procedure.

The variants SakSTAR(K102C) and SakSTAR(K109C), were constructed by the spliced overlap extension polymerase chain reaction (SOE-PCR) (24) using *pMEX.SakSTAR* encoding SakSTAR as template. Two fragments were amplified by PCR (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C), the first one starting from the 5' end (primer 818A) of the *staphylokinase* gene to the region to be mutagenized (forward primer), the second one from this same region (backward primer) to the 3' end of the gene with primer 818D (5' CAAACAGCCAAGCTTCATTCATTCAGC). The forward and backward primers shared an overlap of around 24 bp (for the construction of K102C: TAT GAT AAG AAT TGC AAA AAA GAA GAA (backward) and TTC TTC TTT TTT GCA ATT CTT ATC ATA (forward), for the construction of K109C: AAA AAG AAG AAA CGT GCT CTT TCC CTA (backward) and TAG GGA AAG AGC ACG TTT CTT CTT TTT (forward)). The two purified fragments were then assembled together in a second PCR reaction with the external primers 818A and 818D (30 cycles: 1 sec at 94°C, 1 sec at 50°C, 10 sec at 72°C). The amplified product from this final reaction was purified, digested with EcoRI and HindIII and ligated into the corresponding site of *pMEX.SakSTAR*. For each construction, the sequence of the variant was confirmed by sequencing the entire *SakSTAR* coding region.

The SakSTAR variants were expressed and purified, as described below, from transformed *E. coli* grown in terrific broth (TB) medium (28). A 2 to 4 mL aliquot of an overnight saturated culture in LB medium was used to inoculate a 1 to 2 L culture in terrific broth supplemented with 100 µg/mL ampicillin. The culture was incubated with vigorous aeration and at 30°C. After about 16 hours incubation, IPTG (200 µmol/L) was added to the culture to

induce expression from the *tac* promoter. After 3 hours induction, the cells were pelleted by centrifugation at 4,000 rpm for 20 min, resuspended in 1/10 volume of 0.01 mol/L phosphate buffer pH 6-6.5 and disrupted by sonication at 0°C. The suspension was centrifuged for 20 min at 20,000 rpm and the supernatant was stored at 4°C or at -20°C until purification. The material was purified essentially as described above (Example 2): cleared cell lysates containing the SakSTAR variants were subjected to chromatography on a 1.6 x 5 cm column of SP-Sephadex, followed by chromatography on a 1.6 x 8 cm column of phenyl-Sepharose. The SakSTAR containing fractions, localized by SDS-gel electrophoresis, were pooled for further analysis.

Biochemical analysis

Protein concentrations were determined according to Bradford (29). SDS-PAGE was performed with the Phast System™ (Pharmacia, Uppsala, Sweden) using 10-15% gradient gels and Coomassie Brilliant blue staining, and the specific activities of SakSTAR solutions were determined with a chromogenic substrate assay carried out in microtiter plates (as described in example 2). The specific activity of the different SakSTAR variants are summarized in Table 13.

Mutant SakSTAR(K102C) was essentially monomeric as visualized by SDS-PAGE and Coomassie Brilliant blue staining. Its specific activity was comparable to that of wild-type staphylokinase. In contrast, SakSTAR(K109C) showed a propensity to form dimers (> 60%). This resulted in a markedly increased specific activity in the plasminogen-coupled chromogenic substrate assay (see Table 13). Upon reduction with dithiothreitol (DTT) (20-fold molar excess during 1.5 hour at 37°C) and alkylation with iodoacetamide (100-fold molar excess during 1 hour at 37°C), the K109C dimer is converted into a stable monomer and its resulting specific activity is within the expected range towards wild-type staphylokinase (Table 13). This results confirms that formation of homodimers is the unique determinant for this large increase in specific activity. Dimeric SakSTAR(K109C) was separated from monomeric SakSTAR(K109C) by chromatography on Source S (Pharmacia) (5 x 50mm). Loading buffer was 10 mM phosphate, pH 6.0 and dimeric SakSTAR(K109C)

was eluted by a salt gradient (up to 1 M) in the same buffer. The dimeric SakSTAR(K109C) (>95% pure) containing fractions, localized by SDS-gel electrophoresis, were pooled for further analysis.

Chemical crosslinking of cysteine mutants of SakSTAR with polyethylene glycol

The thiol group of the cysteine mutant SakSTAR(K102C) was targeted for coupling with an activated polyethylene glycol, OPSS-PEG (Shearwater Polymers Europe, Enschede, The Netherlands). OPSS-PEG is a 5 kDa PEG molecule carrying a single activated thiol group at one end that react specifically at slightly alkaline pH with free thiols. Modification of SakSTAR(K102C) was achieved by incubating the molecule (100 μ M) with a three-fold excess of SS-PEG in a 5 mM phosphate, pH 7.9 solution at room temperature. The extent of the reaction was monitored by following the release of 2-thiopyridone from OPSS-PEG at 412 nm. After reaction (about 15 min), the excess of OPSS-PEG was removed by purifying the derivatized SakSTAR(K102C-PEG) on a 1.6 x 5 cm column of SP-Sephadex as described above (see Example 2). The SakSTAR(K102C-PEG) containing fractions, localized by optical density at 280 nm, were pooled for further analysis. SDS-PAGE analysis and Coomassie blue staining confirmed that PEG crosslinking on SakSTAR(K102C) was quantitative. As shown in Table 13, the specific activity of the PEG-derivative was only marginally affected when compared to that of wild-type staphylokinase.

Fibrinolytic properties of SakSTAR variants in human plasma *in vitro*

The fibrinolytic and fibrinogenolytic properties of SakSTAR variants were determined as previously described. Dose- and time-dependent lysis of 125 I -fibrin labeled human plasma clots submerged in human plasma was obtained with four molecules: SakSTAR(K109C) as dimer and as monomer (after reduction and alkylation with iodoacetamide), the monomeric SakSTAR(K102C) and the PEG-derivatized SakSTAR(K102C). Spontaneous clot lysis during the experimental period was $\leq 5\%$ (not shown). Equi-effective concentrations of test compound (causing 50% clot lysis in 2 hrs; C₅₀), determined graphically from plots of clot lysis at 2 hrs versus the concentration of plasminogen activator (not shown), were

comparable to that of SakSTAR, for monomeric SakSTAR(K109C) and SakSTAR(K102C) (Table 13). However, it was observed that the C_{50} for clot lysis by dimeric SakSTAR(K109C) was only 0.12 $\mu\text{g/ml}$, which is approximately three-fold lower than for wild-type staphylokinase. In contrast, a C_{50} of 0.60 $\mu\text{g/ml}$ was measured for SakSTAR(K102C-PEG), which is only two-fold higher than for wild-type staphylokinase. Thus, dimerization of SakSTAR via disulfide bridges or increasing the size of the molecule via PEG-derivatization does not preclude the fibrinolytic activity of staphylokinase. While a PEG-molecule appears to reduce the diffusion and therefore fibrinolytic potency of the derivatized staphylokinase within a fibrin clot, dimerization of staphylokinase results in a synergistic fibrinolytic effect on human fibrin clots.

Pharmacokinetic properties of dimeric SakSTAR(K109C) and SakSTAR(K102C-PEG) following bolus injection in hamsters

The pharmacokinetic parameters of the disposition of dimeric SakSTAR(K109C) and SakSTAR(K102C-PEG) from blood were evaluated in groups of 4 hamsters following intravenous bolus injection of 100 $\mu\text{g/kg}$ SakSTAR variant. SakSTAR-related antigen was assayed using the ELISA described elsewhere. The ELISA was calibrated against each of the SakSTAR variants to be quantitated. Pharmacokinetic parameters included: initial half-life (in min), $t_{1/2\alpha} = \ln 2/\alpha$; terminal half-life (in min), $t_{1/2\beta} = \ln 2/\beta$; volume of the central (plasma) compartment (in mL), $V_C = \text{dose}/(A+B)$; area under the curve (in $\mu\text{g}\cdot\text{min}\cdot\text{mL}^{-1}$), $\text{AUC} = A/\alpha + B/\beta$; and plasma clearance (in $\text{mL}\cdot\text{min}^{-1}$), $\text{Clp} = \text{dose}/\text{AUC}$ (32).

The disposition rate of staphylokinase-related antigen from blood following bolus injection of 100 $\mu\text{g/kg}$ of the selected SakSTAR variants in groups of 4 hamsters could adequately be described by a sum of two exponential terms by graphical curve peeling (results not shown), from which the pharmacokinetic parameters $t_{1/2\alpha}$ and Clp , summarized in Table 13 were derived. The pharmacokinetic parameters of dimeric SakSTAR(K109C) and SakSTAR(K102C-PEG) were markedly different from those of wild type SakSTAR. Initial plasma half-lives ($t_{1/2(\alpha)}$) were 3.6 and min and plasma clearances (Clp) were 0.52 and mL/min , for dimeric SakSTAR(K109C) and SakSTAR(K102C-PEG), respectively.

06/27/2019

CONCLUSION

In summary, the present experience illustrates that staphylokinase variants with markedly *reduced* antibody induction but *intact* thrombolytic potency can be generated. To our knowledge, this observation constitutes the first case in which a heterologous protein, with the use of protein engineering techniques, is rendered significantly less immunogenic in man without reducing its biological activity.

The present invention was initiated by the observation that certain "clustered charge-to-alanine" substitution variants of recombinant staphylokinase (SakSTAR variant (9)) had a reduced reactivity with antibodies induced by treatment with wild type SakSTAR (3,4) and induced less antibodies than wild type SakSTAR in patients with peripheral arterial occlusion (22,32,35). In an effort to optimize the specific activity versus antigenicity ratio, a comprehensive mutagenesis study, comprising the construction and expression of over 250 plasmids encoding SakSTAR variants, and the purification of the translation products was undertaken. The SakSTAR variants were characterized in terms of specific activity, affinity towards a panel of murine monoclonal antibodies and absorption of SakSTAR specific antibodies from pooled plasma of 10 patients treated with wild type SakSTAR and of two subpools of 3 patients each which reacted strongly (subpool B) or poorly (subpool C) with the immunodominant epitope K74,E75,R77. In a later phase, an additional pool of 40 patients treated with wild-type SakSTAR was also used for absorption studies. The values obtained with both pools were in good agreement. Linear regression analysis yielded: (Pool 10)= (Pool 40) +, with $r = \dots$

Residues for site-directed mutagenesis were selected in three ways: 1) a comprehensive analysis of variants with 1 or 2 adjacent amino acids substituted with Ala; 2) analysis of the differential reactivity of the two natural variants SakSTAR and Sak42D (which corresponds to SakSTAR(S34G,G36R,H43R) and 3) surface exposure of the residues as derived from the three dimensional structure. From these analyses, SakSTAR(K35A), SakSTAR(N95A) and SakSTAR(S103A) emerged with specific activities ≥ 200 kU/mg, SakSTAR(W66A), SakSTAR(Y73A) and SakSTAR(E75A) with reduced reactivity with ≥ 3 of the 5 antibodies

of epitope cluster I, SakSTAR(H43A) and SakSTAR(V45A) with 3 antibodies of epitope cluster II, and SakSTAR(V32A), SakSTAR(K35A), SakSTAR(D82A) and SakSTAR(K130A) and ≥ 3 of the 5 antibodies of epitope cluster III. However, only SakSTAR(Y73A) and SakSTAR(K74A) reduced the antibody binding from pooled immunized patient plasma by ≥ 30 percent. Analysis of the differential reactivity of SakSTAR and Sak42D revealed that the reduced reactivity with antibodies of epitope cluster III and with immunized patient plasma could be ascribed to the G36R substitution. In addition E65 and K135 of SakSTAR were targeted because of their location, in the threedimensional structure (36), in the close vicinity of the immunodominant K74,E75,R77 epitope.

Substitution of K35 with other amino acids than Ala did not improve the specific activity over that of SakSTAR(K35A), and substitution of Y73 with other residues did not rescue the impaired specific activity. SakSTAR(E80A,D82A) had an intact specific activity and a somewhat lower reactivity with immunized patient plasma, and the increased specific activity of SakSTAR(N95A) could not be conferred to variants containing K74A or K135R. SakSTAR(K130A) was outperformed in terms of specific activity by SakSTAR(K130T). K74 confirmed its key role in binding of antibodies from immunized patient plasma in the absence of a markedly reduced reactivity with the murine monoclonal antibodies (22), the best specific activity/antigenicity ratios being obtained with SakSTAR(K74Q) and SakSTAR(K74R).

The SakSTAR(K130T,K135R) variant was taken as a template for additive mutagenesis because of its high specific activity with a moderate reduction of binding to antibodies of epitope cluster III and of absorption of antibodies from immunized patient plasma (Table 6). Addition of G36R, K74R or both to the template did not affect the specific activity, but reduced the reactivity with monoclonal antibodies against epitope cluster III (G36R substitution) and decreased the absorption of antibodies from immunized patient plasma (K74R substitution). Addition of E80A and/or D82A to the SakSTAR(K130T,K135R) template did not affect the specific activity and was selected as a template for further mutagenesis because of its reduced reactivity with epitope cluster III antibodies (Table 7). Addition of K74R and even more of K74Q drastically reduced the reactivity with immunized

patient plasma. Finally, addition of E65D or E65Q to the SakSTAR(K74R,E80A,D82A,K130T,K135R) template yielded variants with intact specific activity which only bound 1/3 of the antibodies of pooled immunized patient plasma, only about 10 to 30 percent of the antibodies from plasma of patients with a high concentration of antibodies directed towards the immunodominant K74,E75,R77 (Subpool B) and only about 60 percent of the antibodies from plasma of patients with a very low concentration of antibodies directed against this immunodominant epitope (Subpool C). Based on this analysis, SakSTAR(K74Q,E80A,D82A,K130T,K135R), and SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) were selected for further analysis.

The fibrinolytic potency and the fibrin-selectivity of these selected mutants in a plasma milieu was indistinguishable from that of wild type SakSTAR. The temperature stability of the mutants was still acceptable with no significant loss of activity upon incubation at 37°C for 3 days, although at 56° and 70°C, they were more rapidly inactivated than wild-type SakSTAR. The pharmacokinetics of the SakSTAR variants following intravenous bolus injection in hamsters did not reveal major differences with wild type SakSTAR except for a possibly somewhat higher plasma clearance.

In conclusion, the two variants of SakSTAR which emerged from the present site directed mutagenesis program are characterized by an intact or slightly increased specific activity, maintained thrombolytic potency and fibrin-selectivity in a human plasma milieu, acceptable although slightly reduced temperature stability and a markedly reduced reactivity with anti-SakSTAR antibodies in pooled immunized patient plasma. In view of the previously found correlation between reduced antigenicity and reduced immunogenicity of certain "charged-cluster-to-alanine" variants investigation of immunogenicity associated with their use for thrombolytic therapy in man appeared warranted.

Highly purified, sterilized preparations of SakSTAR(K74Q,E80A,D82A,K130T,K135R) and SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) were produced and found to contain low endotoxin levels, and to be devoid of acute toxicity in mice following intravenous bolus injection at a dose of 3 mg/kg.

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REFERENCES

13. Lijnen HR, Van Hoef B, Matsuo O, Collen D: On the molecular interactions between plasminogen-staphylokinase, α_2 -antiplasmin and fibrin. *Biochim Biophys Acta* 1118: 144-148, 1992.
14. Silence K, Collen D, Lijnen HR: Interaction between staphylokinase, plasmin(ogen) and α_2 -antiplasmin. Recycling of staphylokinase after neutralization of the plasmin-staphylokinase complex by α_2 -antiplasmin. *J Biol Chem* 268: 9811-9816, 1993.
15. Silence K, Collen D, Lijnen HR: Regulation by α_2 -antiplasmin and fibrin of the activation of plasminogen with recombinant staphylokinase in plasma. *Blood* 82: 1175-1183, 1993.
16. Sakharov DV, Lijnen HR, Rijken DC. Interactions between staphylokinase, plasmin(ogen), and fibrin. *J Biol Chem* 271: 27912-27918, 1996.
17. Schlott B, Gührs KH, Hartmann M, Röcker A, Collen D. Staphylokinase requires NH_2 -terminal proteolysis for plasminogen activation. *J Biol Chem* (in press).
18. Collen D, De Cock F, Vanlinthout I, Declerck PJ, Lijnen HR, Stassen JM. Comparative thrombolytic and immunogenic properties of staphylokinase and streptokinase. *Fibrinolysis* 6: 232-242, 1992.
19. Collen D, De Cock F, Stassen JM. Comparative immunogenicity and thrombolytic properties toward arterial and venous thrombi of streptokinase and recombinant staphylokinase in baboons. *Circulation* 87: 996-1006, 1993.
20. White H: Thrombolytic treatment for recurrent myocardial infarction. *Br Med J* 302: 429-430, 1991.
21. Gase A, Hartmann M, Gührs KH, Röcker A, Collen D, Behnke D, Schlott B: Functional significance of NH_2 - and COOH -terminal regions of staphylokinase in plasminogen activation. *Thromb Haemost* 76: 755-760, 1996.
22. EP 95200023.0 (January 6, 1995) and US 08/499,092 (July 6, 1995).
23. Schlott B, Hartmann M, Gührs KH, Birch-Hirschfeld E, Pohl HD, Vanderschueren S, Van de Werf F, Michoel A, Collen D, Behnke D: High yield production and purification

- of recombinant staphylokinase for thrombolytic therapy. *Bio/technology* 12: 185-189, 1994.
24. Horton RM, Hunt HD, Ho SN, Pullen JK, Pease LR. Engineering hybrid genes without the use of restriction enzymes: gene splicing by overlap extension. *Gene* 77: 61-68, 1989.
 25. BIAcore system manual, 5-2, Pharmacia Biosensor AB, Uppsala, Sweden.
 26. Karlsson R, Michaelsson A, Mattsson L: Kinetic analysis of monoclonal antibody-antigen interactions with a new biosensor based analytical system. *J Immunol Methods* 145: 229-240, 1991.
 27. Sambrook J, Fritsch EF, Maniatis T: *Molecular cloning: a laboratory manual*. 2nd Ed. Cold Spring Harbor, NY. Cold Spring Harbor Laboratory Press, 1989.
 28. Tartof KD, Hobbs CA: Improved media for growing plasmid and cosmid clones. *Bethesda Res Lab Focus* 9: 12, 1987
 29. Bradford MM: A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. *Anal Biochem* 72: 248, 1976.
 30. Deutsch DG, Mertz ET: Plasminogen: purification from human plasma by affinity chromatography. *Science* 170: 1095-1096, 1970.
 31. Collen D, Moreau H, Stockx L, Vanderschueren S. Recombinant staphylokinase variant with altered immunoreactivity. II. Thrombolytic properties and antibody induction. *Circulation* 94: 207-216, 1996.
 32. Vanderschueren S, Stockx L, Wilms G, Lacroix H, Verhaeghe R, Vermeylen J, Collen D: Thrombolytic therapy of peripheral arterial occlusion with recombinant staphylokinase. *Circulation* 92: 2050-2057, 1995.
 33. Gibaldi M, Perrier D. *Pharmacokinetics*, Marcel Dekker, New York, N.Y., 1983, 45-111.
 34. Inada Y, Furukawa M, Sasaki H, Koderia Y, Hiroto M, Nishimura H, Matsushima A. Biomedical and biotechnological applications of PEG- and PM-modified proteins, *TIBTECH* 13: 86-91, 1995.

35. Collen D, Bernaerts R, Declerck P, De Cock F, Demarsin E, Jenné S, Laroche Y, Lijnen HR, Silence K, Verstreken M. Recombinant staphylokinase variants with altered immunoreactivity. I. Construction and characterization. *Circulation* 94: 197-206, 1996.
36. Rabijns A, De Bondt HL, De Ranter C. Three-dimensional structure of staphylokinase, a plasminogen activator with therapeutic potential. *Nature Struct Biol* 4: 357-360, 1997.

1. The first part of the paper is devoted to the study of the asymptotic behavior of the solutions of the system (1) as $\epsilon \rightarrow 0$. It is shown that the solutions of the system (1) converge to the solutions of the system (2) as $\epsilon \rightarrow 0$.

Table 1: Alanine-to-wild-type” reversal variants of “charged-cluster-to-alanine” mutants of SakSTAR: Association constants ($K_A \times 10^7$ mol/L⁻¹) for the binding to insolubilized murine monoclonal antibodies (Mabs), and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs																SakSTAR patient plasma		
			Epitope I				Epitope II				Epitope III										
			17G11	26A2	30A2	2B12	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10	Pool	Subpool B	Subpool C	
SakSTAR		130	22	13	2.9	7.8	11	38	7.4	19	7.7	2.4	4.0	14	5.4	2.9	0.6	95	95	95	
SakSTAR(K35A,E38A)		97	15	22	4.2	11	7.9	110	10	15	12	2.2	<0.1	<0.1	<0.1	1.0	1.0	93	91	94	
SakSTAR(K74A,E75A,R77A)		110	11	<0.1	<0.1	<0.1	<0.1	150	17	28	14	3.3	2.4	11	4.0	2.1	0.9	55	43	95	
SakSTAR(K35A,E38A,K74A,E75A,R77A)		50	11	<0.1	<0.1	<0.1	<0.1	110	36	26	15	2.0	<0.1	<0.1	<0.1	1.5	1.2	52	41	92	
SakSTAR(E38A,K74A,E75A,R77A)		43	11	<0.1	<0.2	<0.1	<0.1	140	39	26	15	2.1	<0.1	3.2	3.7	1.6	1.1	50	44	95	
SakSTAR(K35A,K74A,E75A,R77A)		56	9.2	<0.1	0.15	<0.1	<0.1	52	14	29	8.8	2.3	<0.1	1.8	<0.1	1.8	0.8	46	43	95	
SakSTAR(K35A,E38A,E75A,R77A)		44	11	0.3	0.1	0.2	<0.1	75	9.8	12	7.3	1.6	<0.1	<0.1	<0.1	0.53	0.64	92	87		
SakSTAR(K35A,E38A,K74A,R77A)		41	8.8	2.9	<0.1	2.0	0.33	110	29	31	10	2.0	<0.1	<0.1	<0.1	0.63	0.74	56	50	93	
SakSTAR(K35A,E38A,K74A,E75A)		19	13	<0.1	0.1	<0.1	<0.1	180	41	37	15	1.6	<0.1	<0.1	<0.1	1.2	0.45	48	41	92	
SakSTAR(E38A,E75A,R77A)		88	11	0.6	0.15	0.4	0.3	79	12	15	10	2.0	<0.1	2.6	4.7	1.1	0.81	95	88	95	
SakSTAR(E38A,E75A)		66	16	0.3	<0.1	<0.1	0.9	56	11	13	8.9	2.0	<0.1	20	4.8	1.3	1.6	91	90	95	
SakSTAR(K35A,E75A,R77A)		68	9.2	<0.1	<0.1	<0.1	<0.1	60	7.0	13	11	3.3	<0.1	1.5	<0.1	0.8	1.1	88	89	95	
SakSTAR(K35A,E75A)		150	17	0.12	<0.1	0.16	0.14	40	7.2	13	9.2	4.2	<0.1	1.8	<0.1	1.4	1.5	94	93	95	
SakSTAR(K74A)		100	12	7.6	0.17	4.4	2.1	55	15	33	14	3.6	2.9	14	4.9	3.4	1.2	59	45	95	
SakSTAR(E75A)		140	13	1.2	<0.1	<0.1	<0.1	46	8.5	14	12	3.4	4.5	18	5.0	1.2	2.1	95	93	95	
SakSTAR(K74A,E75A,R77A,E80A,D82A)		50	14	<0.1	<0.1	<0.1	<0.1	180	19	33	19	3.7	<0.1	<0.1	<0.1	<0.1	1.2	49	29	89	
SakSTAR(E80A,D82A)		130	7.3	12	2.1	6.5	5.9	79	6.1	8.4	7.8	1.9	<0.1	<0.1	<0.1	<0.1	0.44	89	83	92	
SakSTAR(E80A)		160	13	13	3.3	7.9	10	35	7.4	17	8.6	2.1	<0.1	16	3.6	<0.1	1.7	94	93	95	
SakSTAR(D82A)		160	17	12	4.8	7.3	11	31	7.8	17	12	2.7	<0.1	0.18	<0.1	<0.1	2.3	95	93		
SakSTAR(E75A,D82A)		170	20	15	3.1	6.6	7.2	69	8.1	15	14	4.9	0.17	0.7	0.5	0.1	1.4	95	95		

Apparent association constants ≥ 10 -fold lower than those of wild-type SakSTAR are represented in bold type; Spec. Act. $\geq 100,000$ HU/mg represented in bold type; $\leq 60\%$ absorption represented in bold type.

Table 2: Baseline characteristics and treatment outcome of the patients with peripheral arterial occlusion treated with SakSTAR, SakSTAR(K74A) or SakSTAR(K74A,E75A,R77A)

Compound Patient Id.	Gender	Age (yrs)	Clinical ischemia	Locus of occlusion	Age of occlusion (days)	Length of occlusion (cm)	Recanalization by thrombolysis	Total dose of thrombolytic agent (mg)	Total duration of infusion (hrs)	Additional therapy
SakSTAR										
MEE	F	67	Rest pain	Left SFA	30	8	complete	7.0	5.0	PTA
FOR	M	68	Claudication	Left IA (stent)	14	18	complete	6.5	4.5	PTA + stent
DAN	M	73	Claudication	Right SFA	30	6	complete	7.5	5.5	PTA
BER	F	63	Rest pain	Left FT graft	18	55	complete	18	28	PTA
DAM	F	43	Acute	Left brachial and radial artery	2	7	complete	19	17	PTA + stent
TOR	M	68	Claudication	Right SFA (popliteal aneurysm)	50	12	complete	6.0	4.0	PTA + femoropopliteal bypass graft
CLA	M	74	Acute	Left PA	1.5	20	complete	9.0	7.0	-
MAN	M	65	Acute	Left EIA (stent)	4	20	complete	6.5	4.5	(amputation left digit V)
MAT	M	64	Subacute	Right FP graft	3	45	complete	8.0	6.0	(-)
Mean ± SEM		65 ± 3.0			17 ± 5.6	21 ± 5.8		9.7 ± 1.7	9.1 ± 2.7	
SakSTAR(K74A)										
LJE	M	70	Subacute	Right FF graft	10	48	complete	11	9.0	PTA
ENG	M	50	Claudication	Right SFA	28	10	complete	12	10	PTA
COX	F	48	Claudication	Right PA graft	25	7	partial	15	15	PTA
MAN	F	68	Claudication	Right SFA	≥120	9	complete	9.0	7.0	PTA
VHE	M	47	Acute	Right IF graft	10	54	complete	18	16	Surgical graft revision
MUL	F	51	Acute	Right IF and FP graft	1	63	complete	16	20	PTA
BUR	F	67	Rest pain	Right TF trunc	9.0	38	partial	18	21	-
NIJ	F	60	Rest pain	Left AF graft	23	78	complete	15	21	-
POE*	M	49	Subacute	Right TF trunc	2	30	partial	6.0	4.0	rt-PA, surgical graft lengthening
VBE	M	39	Subacute	Right BA (embolism)	20	28	complete	18	23	Stent right SC artery, first rib resection
SME	F	50	Subacute	TF trunc	18	32	complete	21	19	None
WOL	M	67	Subacute	Right PA	4	25	complete	16	22	-
Mean ± SEM		56 ± 3.0			23 ± 9.2	35 ± 6.4		15 ± 1.2	16 ± 1.9	
SakSTAR(K74A,E75A,R77A)										
JAC	F	65	Acute	Right BA and UA	0.3	5	complete	14	12	-
MAE	M	74	Rest pain	Left SFA	10	50	complete	9.0	7.0	PTA
CRA	F	52	Claudication	Right IA and FA artery	14	28	complete	25	23	PTA + stent
VDB	M	68	Claudication	Left SFA	90	12	complete	9.0	7.0	PTA
DUN	M	71	Subacute	Left SFA	14	6	complete	9.0	7.0	PTA
DEL	M	59	Acute	Right FT graft	3	42	complete	9.0	7.0	PTA
Mean ± SEM		65±3.3			22±14	24±7.8		13 ± 2.6	11 ± 2.6	

AF: aortofemoral; BA: brachial artery; CIA: common iliac artery; FF: femorofibular; FP: femoropopliteal; FI: iliofemoral; IA: iliac artery; IF: iliofemoral; PA: popliteal artery; PTA: percutaneous transluminal angioplasty; SFA: superficial femoral artery; TF: tibiofibular; UA: ulnar artery.

*Previous treatment with SakSTAR in 1994

Table 3: Alanine-substitution variants of SakSTAR: Association constants ($K_A \times 10^7 \text{ mol/L}^{-1}$) for binding to insolubilized murine monoclonal antibodies (Mab) and absorption (percent) of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs															SakSTAR patient plasma		
			Epitope cluster I					Epitope cluster II					Epitope cluster III					Pool	Subpool B	Subpool C
			17G11	26A2	30A2	28I2	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10			
SakSTAR		120	9.3	13	2.9	7.8	11	38	7.4	19	7.7	2.4	4.0	14	5.4	2.9	0.6	95	95	95
			[6.4	17	7.4	19	35	18	25	>18	>14	1.1	0.6	11	16	6.3	2.0]			
SakSTAR(S34G,H43R)		120	10	14	3.3	7.5	11	<0.1	<0.1	<0.1	20	2.7	<0.1	<0.1	<0.1	0.15	1.7	87	76	75
SakSTAR(F4A)	LE																			
SakSTAR(D5A,K6A)		150	[1.3	21	12	9.2	9.7	12	23	17	10	0.4	1.3	3.9	0.8	4.5	3.8]	95	95	95
SakSTAR(K8A,K10A)		24	[1.8	16	5.1	29	15	22	16	26	18	1.0	0.93	11	1.1	18	0.75]	95	95	95
SakSTAR(Y9A)	24	78	22	49	6.6	2.3	16	44	8.4	20	14	2.6	2.4	16	9.6	1.1	0.5	96	95	95
SakSTAR(K11A,D13A,D14A)	LE																			
SakSTAR(D13A)	6	46	2.4	6.1	2.0	3.7	3.4	11	1.9	4.5	4.4	8.7	1.5	2.4	1.1	3.6	<0.1	95	94	95
SakSTAR(D14A)	14	30	5.1	13	4.0	6.6	3.1	38	7.7	12	15	2.2	2.7	6.0	3.2	5.6	<0.1	95	94	95
SakSTAR(S16A)	41	160	8.1	16	4.5	8.4	9.0	15	6.1	13	21	3.6	4.0	8.0	4.3	2.5	0.5	95	95	95
SakSTAR(Y17A,F18A)	27	30	13	22	3.3	10	9.5	21	4.6	6.7	12	2.5	1.2	18	6.5	3.4	<0.1	95	95	95
SakSTAR(E19A,P20A)	36	9	11	19	3.3	9.2	12	15	6.1	11	16	1.0	1.1	15	5.1	3.3	<0.1	93	93	95
SakSTAR(T21A)	54	170	4.8	15	2.4	8.7	9.6	32	11	24	18	1.3	1.8	9.6	2.9	5.6	0.6	95	95	95
SakSTAR(P23A)	41	67	14	31	4.4	14	22	41	5.3	37	13	10	4.0	11	7.6	3.1	1.9	91	95	95
SakSTAR(Y24A)	10	40	17	33	4.3	13	11	33	4.3	7.0	12	4.0	0.4	14	6.8	4.8	<0.1	95	95	95
SakSTAR(L25A)	LE																			
SakSTAR(M26A)	LE																			
SakSTAR(V27A)	62	50	3.3	15	1.6	7.8	7.4	12	2.9	3.7	33	2.0	1.3	5.9	2.8	4.2	1.2	95	95	95
SakSTAR(N28A)	90	<5	5.8	29	2.1	7.0	5.5	27	10	20	2.5	2.1	2.1	5.6	2.1	2.1	0.7	95	95	95
SakSTAR(N28A,V29A)	32	45	18	30	2.5	20	18	20	14	20	24	2.7	3.3	20	1.1	5.0	2.0	93	95	95
SakSTAR(T30A)	52	140	7.4	13	2.1	7.0	6.1	7.6	3.4	5.1	13	3.3	5.6	12	3.4	5.4	0.8	94	95	95
SakSTAR(V32A)	78	45	10	9.6	2.4	6.2	7.8	56	17	12	14	1.4	<0.1	<0.1	<0.1	<0.1	2.2	90	93	95
SakSTAR(D33A,K35A)		130	[2.1	19	14	14	19	15	24	32	10	5.3	1.4	5.1	3.8	5.1	3.0]	95	95	95
SakSTAR(S34A)	29	110	17	24	4.6	9.5	11	28	11	22	15	2.9	3.1	8.8	3.8	2.0	0.2	95	95	93

Table 3 - cont'd: Alanine-substitution variants of SakSTAR: Association constants ($K_A \times 10^7 \text{ mol/L}^{-1}$) for binding to insolubilized murine monoclonal antibodies (Mab) and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs																	SakSTAR patient plasma		
			Epitope cluster I					Epitope cluster II					Epitope cluster III							Pool	Subpool B	Subpool C
			17G11	26A2	30A2	2B12	3G10	18H12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10					
SakSTAR(K35A)		230	6.4	14	3.3	8.0	7.4	31	11	12	11	2.6	<0.1	1.3	0.2	1.7	0.8	91	88	95		
SakSTAR(K35A,E38A)		97	15	22	4.2	11	7.9	110	10	15	12	2.2	<0.1	<0.1	<0.1	1.0	1.0	93	91	94		
SakSTAR(G36A)	14	72	3.5	9.8	1.5	5.7	6.5	52	4.2	17	9.2	1.4	<0.1	<0.1	0.9	5.0	1.0	86	83	78		
SakSTAR(N37A)	40	110	5.6	31	3.0	10	11	20	4.1	14	10	2.9	1.4	5.3	3.5	3.6	0.8	95	95	95		
SakSTAR(L39A,L40A)	8	<5	14	15	3.1	5.1	8.0	27	16	6.3	12	2.7	1.2	5.4	3.2	2.1	0.9	93	93	95		
SakSTAR(S41A,P42A)	24	48	10	25	4.1	13	12	11	3.0	1.9	27	2.7	3.2	15	4.8	3.6	1.1	95	95	95		
SakSTAR(H43A)	37	69	15	28	9.7	18	7.6	<0.1	<0.1	<0.1	9.1	1.5	2.0	23	7.8	7.2	1.6	95	95	95		
SakSTAR(H43A,Y44A)	13	<5	10	22	3.7	17	15	<0.1	<0.1	<0.1	1.4	3.0	2.3	11	5.2	2.1	0.1	95	95	95		
SakSTAR(V45A)	19	<5	16	5.6	1.4	4.8	6.3	2.2	0.2	1.7	32	2.6	2.1	8.3	1.1	2.8	1.6	91	92	95		
SakSTAR(E46A,K50A)	LE																					
SakSTAR(F47A)	6	<5	<0.1	4.0	1.0	3.9	3.4	5.7	2.7	2.8	8.5	1.7	0.9	8.8	3.0	3.0	0.9	90	82	97		
SakSTAR(I49A)	2	43	2.7	27	7.8	23	22	35	4.4	11	6.2	1.1	2.0	5.7	2.0	1.7	0.6	95	95	95		
SakSTAR(K50A)	15	42	<0.1	13	2.9	7.8	8.7	46	8.3	12	2.2	0.5	2.8	6.4	4.0	2.3	0.6	95	94	95		
SakSTAR(T53A,T54A)	14	68	0.9	19	2.7	7.6	7.8	41	6.7	12	15	1.5	1.9	5.1	2.3	1.0	0.6	93	94	95		
SakSTAR(L55A)	LE																					
SakSTAR(T56A)	17	150	5.5	15	3.2	12	13	56	5.3	12	11	2.0	3.5	6.1	2.7	4.3	1.2	94	92	95		
SakSTAR(K57A,E58A,K59A)		94	[4	8.7	6.0	7.3	27	16	14	6.7	5.6	0.52	0.36	1.7	0.42	1.0	1.1]					
SakSTAR(I60A)	11	96	12	20	2.9	11	13	27	6.4	25	2.7	1.5	0.7	5.8	2.9	1.7	1.0	95	95	95		
SakSTAR(E61A,E65A)		80	[9.5	>10	8.8	21	29	>11	>16	6.6	>7.2	4.6	0.5	4.6	2.0	5.9	1.5]					
SakSTAR(Y62A,Y63A)	24	<5	<0.1	4.3	0.3	2.1	1.9	11	2.2	3.1	8.4	1.7	0.6	9.2	3.6	3.8	0.7	89	83	95		
SakSTAR(Y63A)	4	<5	<0.1	18	3.7	9.6	13	17	5.3	3.3	15	1.1	2.2	5.3	4.3	1.0	3.7	89	82	95		
SakSTAR(Y64A)	14	48	14	16	2.9	6.3	7.8	15	15	21	21	2.6	1.6	7.6	5.6	2.8	0.7	94	92	95		
SakSTAR(E65A)	25	97	53	20	4.4	12	7.0	10	5.6	9.7	1.9	1.8	2.3	4.7	3.0	5.8	0.97	95	95	95		
SakSTAR(E65A,D69A)		<5																				
SakSTAR(W66A)	26	<5	<0.1	<0.1	<0.1	<0.1	<0.1	15	4.4	5.7	23	3.3	2.0	10	1.4	1.8	0.8	85	78	92		
SakSTAR(L68A)	16	93	14	22	3.5	8.5	9.3	29	8.7	16	15	4.0	2.1	5.3	3.6	4.1	<0.1	92	92	95		
SakSTAR(T71A)	LE																					

antibodies (Mab) and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs																SakSTAR patient plasma			
			Epitope cluster I						Epitope cluster II						Epitope cluster III				Pool	Subpool B	Subpool C	
			17G11	26A2	30A2	2B12	3G10	18P12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10					
SakSTAR(Y73A)	20	<5	9.5	<0.1	<0.1	<0.1	4.8	33	7.3	11	8.9	2.0	0.5	9.0	2.5	1.4	0.8	63	44	93		
SakSTAR(Y73A,K74A)	24	<5	18	<0.1	<0.1	<0.1	19	6.7	23	9.9	3.2	2.7	13	4.0	1.6	1.1	47	28	87			
SakSTAR(K74A)	80	69	4.4	2.7	0.2	2.2	1.1	17	5.2	14	7.6	2.2	2.0	6.8	3.3	1.8	0.9	64	58	95		
SakSTAR(K74A,E75A,R77A)		68	9.2	<0.1	<0.1	<0.1	60	7.0	13	11	3.3	<0.1	1.5	<0.1	0.8	1.1	88	89	95			
SakSTAR(K74A,R77A)	34	41	3.5	1.8	0.2	1.5	0.4	20	2.4	10	2.1	1.8	1.7	2.3	2.2	1.2	0.7	74	60	95		
SakSTAR(E75A)		140	13	1.2	<0.1	<0.1	46	8.5	14	12	3.4	4.5	18	5.0	1.2	2.1	95	93	95			
SakSTAR(F76A)	9	90	20	9.6	1.0	2.7	3.9	13	6.2	20	1.5	1.7	0.3	5.9	2.1	1.2	1.0	94	92	95		
SakSTAR(V78A,V79A)	23	68	12	23	4.0	10	17	21	18	34	28	2.3	1.6	4.7	<0.1	0.5	1.7	93	93	95		
SakSTAR(E80A)		160	13	13	3.3	7.9	10	35	7.4	17	8.6	2.1	<0.1	16	3.6	<0.1	1.7	94	93	95		
SakSTAR(E80A,D82A)		130	7.3	12	2.1	6.5	5.9	79	6.1	8.4	7.8	1.9	<0.1	<0.1	<0.1	0.4	89	83	92			
SakSTAR(L81A)	23	28	12	33	1.6	40	11	52	11	17	17	3.9	1.4	5.2	7.1	4.6	1.5	88	95	95		
SakSTAR(D82A)		160	17	12	4.8	7.3	11	31	7.8	17	12	2.7	<0.1	0.2	<0.1	2.3	95	93	95			
SakSTAR(D82A,S84A)	72	130	8.3	14	2.6	8.1	8.5	23	3.8	12	11	1.7	<0.1	<0.1	1.4	<0.1	1.0	91	91	95		
SakSTAR(S84A)	12/26	89	8.0	16	3.8	8.6	10	90	8.3	11	3.6	1.8	2.2	1.6	3.0	3.5	0.5	95	95	95		
SakSTAR(K86A,E88A)		73	7.2	1.4	3.7	6.0	4.6	5.7	4.9	7.7	15	4.4	<0.1	5.4	0.80	1.9	0.13	95	95	95		
SakSTAR(I87A)	18	98	6.7	23	2.8	8.6	9.1	10	3.6	11	7.4	2.7	1.1	7.8	3.4	4.5	1.0	95	95	95		
SakSTAR(V89A)	20	87	4.6	11	2.6	6.6	2.2	28	7.2	7.3	3.0	1.3	1.2	5.1	2.9	3.1	0.83	95	95	95		
SakSTAR(T90A)	78	120	6.0	12	0.9	3.7	3.1	20	4.8	7.2	<0.1	<0.1	2.1	6.6	2.6	2.1	0.5	95	95	95		
SakSTAR(Y91A)	5	53	6.0	16	3.0	7.0	13	28	8.2	16	0.6	2.1	1.4	3.7	1.6	1.6	0.2	95	95	95		
SakSTAR(Y92A)	16	120	16	23	4.1	13	12	29	7.3	18	<0.1	1.7	4.4	10	3.9	5.9	1.1	95	95	95		
SakSTAR(E93A,K94A)		97	8.2	19	13	30	24	18	11	>10	9.0	0.88	1.4	11	2.4	7.0	2.1]					
SakSTAR(K94A,N95A,K97A)	32	8	NT															95	94	95		
SakSTAR(N95A)	25	260	10	18	4.0	10	11	50	13	14	4.9	2.3	3.7	7.3	4.7	2.9	0.8	95	94	95		
SakSTAR(K96A,K97A,K98A)		47	2.8	41	23	37	90	>16	9.1	19	16	0.41	0.58	17	1.2	13	0.30]					
SakSTAR(E99A)	24	42	7.4	15	4.0	8.4	8.9	22	2.7	4.7	<0.1	<0.1	2.1	6.2	7.3	14	0.8	92	91	92		
SakSTAR(E99A,E100A)	LE																					

Table 3 - cont'd: Alanine-substitution variants of SakSTAR: Association constants ($K_A \times 10^7 \text{ mol/L}^{-1}$) for binding to insolubilized murine monoclonal antibodies (Mab) and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs																SakSTAR patient plasma			
			Epitope cluster I				Epitope cluster II								Epitope cluster III				Pool	Subpool B		Subpool C
			17G11	26A2	30A2	2B12	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10					
SakSTAR(T107A)	23	85	4.6	11	2.1	6.6	7.3	30	2.8	16	0.7	1.0	1.3	5.4	2.4	2.9	0.8	95	95	95	95	
SakSTAR(K102A)	12	89	4.9	12	3.7	6.5	6.3	30	3.1	15	8.2	6.1	0.8	4.2	1.9	10	0.6	95	93	95	95	
SakSTAR(S103A)	67	210	9.0	16	5.0	9.4	9.1	19	5.9	13	13	3.6	3.9	8.3	4.7	2.8	0.9	94	95	95	95	
SakSTAR(F104A)	14	55	5.8	19	4.8	14	27	7.3	5.0	14	4.8	<0.1	0.4	7.6	3.4	1.1	1.3	95	93	95	95	
SakSTAR(I106A)	2	93	2.3	13	3.0	7.4	6.7	5.5	5.2	17	11	1.4	1.8	3.1	1.8	1.2	0.5	95	95	95	95	
SakSTAR(T107A)	32	130	5.2	15	3.4	9.8	10	32	8.7	4.7	14	1.9	3.1	6.3	3.2	5.0	0.8	94	94	95	95	
SakSTAR(E108A,K109A)		170	[1.6	5.1	7.2	19	5.1	28	15	21	21	1.2	0.43	6.9	1.4	10	1.9					
SakSTAR(F111A)	5	49	3.7	16	3.8	13	22	21	8.4	12	3.1	0.8	2.8	2.9	1.5	1.5	0.9	95	95	95	95	
SakSTAR(V112A,V113A)	64	130	4.2	16	3.9	10	12	34	5.8	13	8.0	0.3	1.5	4.3	2.3	3.0	0.8	95	95	95	95	
SakSTAR(D115A,S117A)	80	54	3.3	14	4.1	15	15	17	3.4	19	0.7	<0.1	1.5	4.8	2.6	1.3	0.9	95	95	95	95	
SakSTAR(D115A,E118A,H119A)		32	[2.5	32	3.4	21	8.7	13	9.9	23	9.3	1.2	1.0	24	2.1	9.0	1.8					
SakSTAR(L116A,S117A)	25	<5	4.4	35	3.6	33	42	160	29	220	<0.1	<0.1	0.5	4.1	4.9	3.5	1.6	94	95	95	95	
SakSTAR(H119A,K121A)		130	[8.0	24	11	26	29	25	14	29	12	0.52	1.2	11	2.9	20	1.2					
SakSTAR(I120A)	26	75	23	26	5.1	17	16	30	9.8	25	9.0	6.9	3.0	15	5.1	5.2	1.0	93	95	95	95	
SakSTAR(N122A)	5	19	NT															95	93	95	95	
SakSTAR(F125A)	3	<10	2.8	18	4.7	11	18	17	3.2	6.0	1.9	<0.1	0.3	5.3	2.1	0.9	1.6	93	90	95	95	
SakSTAR(N126V)	11	51	7.6	13	2.0	12	13	30	58	290	9.8	2.5	1.8	8.0	4.2	6.5	0.7	95	95	95	95	
SakSTAR(L127A)	11	54	8.9	6.7	1.8	5.0	6.6	25	4.9	14	8.4	1.5	0.9	1.9	0.9	2.5	1.8	93	94	95	95	
SakSTAR(I128A)	10	20	16	25	4.8	15	14	38	2.6	4.3	8.2	2.9	2.5	2.0	4.2	6.7	0.9	95	93	95	95	
SakSTAR(T129A)	44	190	5.3	15	2.3	14	24	21	13	15	4.2	2.3	0.7	10	3.3	1.3	1.0	95	95	95	95	
SakSTAR(K130A)	130	280	5.1	12	3.2	6.4	3.5	22	6.7	11	15	2.7	<0.1	<0.1	4.1	0.9	0.6	92	74	71	71	
SakSTAR(V131A)	130	70	6.5	17	2.9	11	13	19	14	19	29	3.4	1.9	13	5.3	8.6	0.9	95	95	95	95	
SakSTAR(V132A)	100	130	4.2	15	2.6	9.2	11	33	12	30	19	2.1	2.1	3.6	<0.1	2.6	0.4	95	95	95	95	
SakSTAR(I133A)	3	99	9.4	15	1.9	7.8	7.8	24	6.0	9.1	8.6	1.4	0.56	6.4	1.6	1.6	0.9	95	95	95	95	
SakSTAR(E134A,K135A,K136A)		74	[22	21	6.7	25	25	>18	>25	>15	>12	1.7	0.2	11	0.94	6.0	2.6]					
SakSTAR(K135A)	54	410	5.2	12	11	7.9	11	20	11	11	3.8	2.0	1.6	6.9	3.7	1.9	0.9	95	95	95	95	
SakSTAR(K136A)		180	7.6	18	5.6	12	13	54	5.3	12	16	3.2	3.5	8.6	3.9	1.8	0.5	91	83	83	95	

LE: expression level below 3 mg/L

Table 4: Mutagenesis of S34, G36 and H43: Association constants ($K_A \times 10^7 \text{ mol/L}^{-1}$) for binding to insolubilized murine monoclonal antibodies (Mab) and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. Act. (μ U/mg)	murine MAbs																SakSTAR patient plasma		
		Epitope cluster I				Epitope cluster II				Epitope cluster III								Pool	Subpool B	Subpool C
		17G11	26A2	30A2	2B12	3G10	18F12	14H5	28H4	32B2	7F10	7H11	23E1	40C8	24C4	1A10				
SakSTAR		120	9.3	13	2.9	7.8	11	38	7.4	19	7.7	2.4	4.0	14	5.4	2.9	0.6	95	95	95
SakSTAR(S34G,G36R,H43R)		120	10	14	3.3	7.5	11	<0.1	<0.1	<0.1	20	2.7	<0.1	<0.1	<0.1	0.15	1.7	87	76	75
SakSTAR(S34A)	29	110	17	24	4.6	9.5	11	28	11	22	15	2.9	3.1	8.8	3.8	2.0	0.2	95	95	93
SakSTAR(G36A)	14	72	3.5	9.8	1.5	5.7	6.5	52	4.2	17	9.2	1.4	<0.1	<0.1	0.9	5.0	1.0	86	83	78
SakSTAR(G36E)	12	66	3.1	8.7	1.4	4.8	4.7	12	2.8	6.1	7.6	1.0	<0.1	<0.1	<0.1	3.4	1.1	89	83	72
SakSTAR(G36K)	34	88	9.9	23	3.1	8.3	9.8	21	3.9	13	15	3.0	<0.1	<0.1	<0.1	2.6	1.2	88	80	69
SakSTAR(G36L)	15	92	3.6	11	1.8	6.1	6.1	16	1.4	6.4	12	1.3	<0.1	<0.1	<0.1	0.6	1.1	93	85	72
SakSTAR(G36N)	10	91	8.7	10	1.6	6.1	6.2	33	3.3	7.8	7.9	1.8	<0.1	<0.1	1.0	0.3	0.5	86	80	75
SakSTAR(G36Q)	21	92	10	12	1.8	6.7	6.5	23	3.8	7.5	7.3	1.5	<0.1	<0.1	<0.1	0.1	0.4	87	84	73
SakSTAR(G36R)	45	100	11	2.4	3.3	10	10	27	4.6	14	20	3.4	<0.1	<0.1	<0.1	3.1	1.2	89	81	70
SakSTAR(H43A)	37	69	15	28	9.7	18	7.6	<0.1	<0.1	<0.1	9.1	1.5	2.0	23	7.8	7.2	1.6	95	95	95
SakSTAR(H43R)	43	120	15	11	2.7	7.6	11	<0.1	<0.1	<0.1	13	6.4	0.7	18	6.7	5.7	1.4	95	95	95
SakSTAR(S34G,G36R)	45	90	3.1	12	2.3	4.8	4.2	13	8.3	24	9.1	1.9	<0.1	<0.1	<0.1	<0.1	0.6	92	83	69
SakSTAR(S34G,G36R,H43R,K74A)	1	12	12	4.7	3.8	7.4	9.4	<0.1	0.1	0.6	25	2.3	<0.1	<0.1	0.5	0.8	1.7	67	56	83
SakSTAR(S34G,G36R,K74A)	15	26	4.0	2.1	<0.1	1.8	0.6	10	2.2	15	13	1.8	<0.1	<0.1	<0.1	0.3	2.2	59	28	68
SakSTAR(K35G,G36R,H43D)	32	6	1.8	1.5	1.6	2.0	8.9	<0.1	<0.1	<0.1	7.1	1.3	<0.1	<0.1	<0.1	<0.1	0.9	82	75	72
SakSTAR(G36R,K74A)	40	35	19	7.0	0.2	4.3	2.0	53	27	28	19	4.4	<0.1	<0.1	<0.1	1.2	1.0	48	33	58
SakSTAR(G36R,K74R)	68	150	4.7	17	3.8	11	8.0	16	6.0	6.4	3.0	1.6	<0.1	<0.1	<0.1	0.2	0.8	81	54	73
SakSTAR(G36R,K74A,N95A)	11	25	6.1	2.9	2.9	2.4	<0.1	31	5.7	12	5.3	4.1	<0.1	<0.1	<0.1	<0.1	0.9	53	32	63
SakSTAR(G36R,K74A,K135R)	20	33	5.8	3.4	<0.1	1.7	0.7	26	16	14	17	1.2	<0.1	<0.1	<0.1	0.4	0.5	64	32	68
SakSTAR(G36R,K74R,K135R)	48	75	6.1	17	5.8	10	3.3	31	14	13	5.7	2.3	<0.1	<0.1	<0.1	0.3	0.8	77	49	68

Table 5 - cont'd: Mutagenesis of K35, Y73, K74, E80/D82, N95, K130, V132 and K135: Association constants ($K_A \times 10^7 \text{ mol/L}^{-1}$) for binding to insolubilized murine monoclonal antibodies (Mab) and absorption (percent) of antibodies of immunized patient plasma

Variant	Exp. (mg/L)	Spec. Act. (kU/mg)	murine MAbs														SakSTAR patient plasma	
			Epitope cluster I							Epitope cluster II							Pool	Subpool B
			77G11	26A2	30A2	28B2	30G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4		
SakSTAR(K130A)		280	5.1	12	3.2	6.4	3.5	22	6.7	11	15	2.7	<0.1	<0.1	4.1	0.9	92	74
SakSTAR(K130T)		290	7.8	14	3.1	8.0	5.6	43	5.4	9.7	17	2.7	<0.1	<0.1	3.4	0.5	92	84
SakSTAR(V132A)	102	130	4.2	15	2.6	9.2	11	33	12	30	19	2.1	2.1	3.6	<0.1	2.6	95	95
SakSTAR(V132L)	126	120	4.8	14	2.3	8.0	9.7	67	10	48	20	2.5	2.0	1.1	<0.1	4.8	95	95
SakSTAR(V132T)	78	140	4.5	13	2.4	7.8	9.0	39	10	25	16	2.0	1.1	2.0	<0.1	<0.1	95	95
SakSTAR(V132N)	16	150	4.5	11	1.7	7.0	7.2	21	17	20	15	2.0	1.5	1.3	<0.1	4.2	95	95
SakSTAR(V132R)	76	230	5.4	12	0.8	3.3	3.2	23	5.5	7.8	4.8	1.1	2.1	3.4	<0.1	0.4	95	95
SakSTAR(K135A)	54	410	5.2	12	1.1	7.9	11	20	11	11	3.8	2.0	1.6	6.9	3.7	1.9	95	95
SakSTAR(K135F)	68	64	3.9	6.3	1.0	6.1	4.1	13	4.5	8.4	16	2.1	1.6	5.8	1.9	1.5	95	95
SakSTAR(K135R)	54	230	4.0	14	1.4	9.3	5.0	31	8.1	13	23	1.9	2.4	2.4	2.6	2.1	95	95
SakSTAR(K135A,K74A)	20	130	NT														84	50
SakSTAR(Y73A,K74A)	24	<5	18	<0.1	<0.1	<0.1	<0.1	19	6.7	23	9.9	3.2	2.7	13	4.0	1.6	47	28
SakSTAR(Y73F,K74A)	21	6	17	6.7	<0.1	2.1	<0.1	42	19	21	20	4.9	1.9	2.8	6.6	2.1	51	34
SakSTAR(I160A,K74A,N95A)		84	5.3	2.7	<0.1	2.5	5.3	17	7.2	9.6	3.4	2.3	2.1	6.2	2.7	2.9	60	47
SakSTAR(N95A,K135R)	120	240	4.9	13	1.9	9.0	9.9	18	12	15	2.5	1.5	1.8	5.9	3.6	3.7	95	95
SakSTAR(K130T,K135R)	15	280	3.7	10	1.6	7.2	8.0	13	7.7	4.5	3.7	1.6	<0.1	<0.1	2.4	0.4	89	60

Table 6: Combination mutants of SakSTAR(K130T,K135R) with K35A, G36R, E65X,K74X and selected other amino acids

Variant	Exp. (mg/mL)	Spec. Act. (kU/mg)	murine MAbs										SakSTAR patient plasma				Code					
			Epitope cluster I					Epitope cluster II					Epitope cluster III									
			17G11	26A2	30A2	2B12	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4		1A10	Pool 10	Subpool B	Subpool C	Pool 40
SakSTAR(K130T,K135R)	15	280	3.7	10	1.6	7.2	8.0	13	7.7	4.5	3.7	1.6	<0.1	<0.1	2.4	0.4	0.6	89	60	73	90	SY2
SakSTAR(G36R,K130T,K135R)	26	220	7.2	18	2.8	11	14	17	3.9	8.9	5.1	1.6	<0.1	<0.1	<0.1	<0.1	1.1	79	65	69	-	SY3
SakSTAR(K74R,K130T,K135R)	18	310	7.3	27	5.9	16	18	17	3.7	11	5.1	1.5	<0.1	<0.1	3.2	0.7	0.9	76	49	69	78	SY4
SakSTAR(K74Q,K130T,K135R)	64	190	4.0	7.2	3.0	7.7	0.9	20	5.7	7.2	8.4	1.8	<0.1	<0.1	2.7	0.7	1.0	50	25	67	62	SY41
SakSTAR(G36R,K74R,K130T,K135R)	5	210	7.6	26	5.7	17	19	17	4.7	10	5.0	1.4	<0.1	<0.1	<0.1	<0.1	0.8	73	44	69	75	SY5
SakSTAR(G36R,K74Q,K130T,K135R)	88	120	5.5	7.3	0.8	11	6.4	35	11	7.1	6.1	2.6	<0.1	<0.1	<0.1	<0.1	0.8	51	25	63	54	SY42
SakSTAR(G36R,H43R,K74R,K130T,K135R)	29	160	5.6	8.8	2.2	10	10	<0.1	<0.1	<0.1	6.3	2.0	<0.1	<0.1	<0.1	<0.1	0.9	72	39	69	-	SY9
SakSTAR(S34G,G36R,K74Q,K130T,K135R)	40	76	4.8	5.9	0.5	13	1.5	18	8.7	7.5	5.1	2.1	<0.1	<0.1	<0.1	<0.1	1.1	52	25	65	61	SY43
SakSTAR(E65A,K74Q,K130T,K135R)	46	170	11	9.0	2.3	11	13	19	66	12	5.7	2.3	<0.1	<0.1	3.5	1.2	0.6	45	16	77	55	SY48
SakSTAR(G36R,E65A,K74Q,K130T,K135R)	80	83	4.7	12	1.2	16	21	27	12	10	6.9	2.6	<0.1	<0.1	<0.1	<0.1	0.7	44	18	65	46	SY44
SakSTAR(G36R,E65A,K74A,K130A,K135R)	17	71	5.7	10	1.8	13	12	21	8.4	6.9	4.6	1.8	<0.1	<0.1	<0.1	<0.1	1.0	41	14	64	50	SY59
SakSTAR(E65A,A72S,K74Q,K130T,K135R)	60	96	5.6	6.0	2.0	3.6	4.9	21	8.6	8.8	2.8	2.5	<0.1	<0.1	3.5	1.7	1.1	51	13	66	56	SY51
SakSTAR(E65Q,K74Q,K130T,K135R)	40	150	6.7	18	2.1	15	16	9.0	3.1	4.1	6.3	2.3	<0.1	<0.1	3.8	0.9	0.6	53	29	67	65	SY49
SakSTAR(K74Q,K86A,K130T,K135R)	54	130	2.4	4.9	<0.1	7.4	3.8	19	8.7	7.6	4.7	1.9	<0.1	<0.1	3.5	1.3	1.5	56	32	69	61	SY55
SakSTAR(E65Q,T71S,K74Q,K130T,K135R)	32	210	6.2	13	1.8	10	13	11	3.9	5.0	6.8	23	<0.1	<0.1	3.1	1.1	0.8	49	21	64	59	SY65
SakSTAR(E65Q,K74Q,E75A,K130T,K135R)	36	46	7.7	<0.1	<0.1	<0.1	<0.1	10	3.9	3.2	7.4	2.6	<0.1	<0.1	4.6	1.3	0.6	43	15	62	55	SY66
SakSTAR(E65Q,K74Q,E75D,K130T,K135R)	35	67	7.0	<0.1	<0.1	<0.1	<0.1	13	5.4	4.9	6.6	2.5	<0.1	<0.1	3.2	1.3	1.0	49	29	63	57	SY67
SakSTAR(K74Q,K130T,K135R,K136A,+137A)	19	78	4.3	24	<0.1	2.7	5.6	20	9.6	7.5	5.6	2.2	<0.1	<0.1	1.7	1.1	1.2	37	12	57	50	SY68
SakSTAR(K74Q,K130A,K135R)	28	240	5.6	5.4	0.5	7.5	5.3	21	8.4	14	6.1	2.4	<0.1	<0.1	4.1	2.6	0.7	57	27	78	65	SY56
SakSTAR(E65Q,K74Q,K130A,K135R)	60	230	6.0	14	2.4	15	17	9.0	3.3	5.8	6.3	2.3	<0.1	<0.1	4.3	2.0	0.6	51	32	73	58	SY69
SakSTAR(K74Q,K130E,K135R)	46	300	4.1	4.4	0.8	6.6	3.8	18	8.5	7.8	5.2	2.1	<0.1	<0.1	2.4	0.7	0.9	55	29	64	59	SY57
SakSTAR(E65Q,K74Q,K130A,K135A)	88	170	5.3	8.9	1.7	7.7	11	16	3.4	5.8	6.2	2.4	<0.1	<0.1	3.5	2.4	0.7	55	27	79	55	SY70
SakSTAR(K74Q,K130E,V132R,K135R)	68	170	4.5	4.9	0.4	6.0	5.2	9.0	4.9	4.3	5.6	2.4	<0.1	<0.1	<0.1	<0.1	0.6	51	20	63	56	SY58
SakSTAR(E65Q,K74Q,T90A,K130A,K135R)	36	170	6.2	13	1.8	12	14	7.3	2.5	3.8	<0.1	<0.1	<0.1	<0.1	4.1	1.9	0.5	51	27	69	57	SY71

murine MAbs

Association constants ≥ 10 -fold lower and antibody absorption ≤ 60 percent of wild-type SakSTAR are represented in bold type; $\geq 100,000$ HU/mg represented in bold type. NT: not tested.

Association constants ≥ 10 -fold lower and antibody absorption ≤ 60 percent of wild-type SakSTAR are represented in bold type; $\geq 100,000$ HU/mg represented in bold type. NT: not tested.

Table 7: Combination mutants of SakSTAR(E80A,D82A,K130T,K135R) with K35A, G36R, E65X, K74X, and selected other amino acids

Variant	Exp. (mg/mL)	Spec. Act. (kU/mg)	murine MAbs																SakSTAR patient plasma				Code
			Epitope cluster I					Epitope cluster II					Epitope cluster III						Pool 10	Subpool B	Subpool C	Pool 40	
			17G11	26A2	30A2	28I2	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A1						
SakSTAR(E80A,D82A,K130T,K135R)	20	250	8.4	17	3.1	11	13	18	4.3	11	5.5	1.5	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.0	80	64	68	SY6
SakSTAR(K74R,E80A,D82A,K130T,K135R)	4	220	5.3	31	2.8	18	11	89	8.4	21	5.9	1.8	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.6	74	34	69	SY7
SakSTAR(K74Q,E80A,D82A,K130T,K135R)	27	110	5.1	6.5	1.2	7.9	7.3	28	6.4	19	4.6	1.7	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.8	46	17	60	SY15
SakSTAR(K35A,K74R,E80A,D82A,K130T,K135R)	70	160	5.9	5.7	6.4	2.7	18	19	16	8.2	9.0	1.8	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.8	66	34	66	SY17
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)	22	140	5.4	50	2.9	49	21	31	11	37	3.4	1.8	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.6	43	11	68	SY19
SakSTAR(E65S,K74R,E80A,D82A,K130T,K135R)	3	110	3.2	12	4.6	44	15	19	5.0	15	6.6	1.4	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.4	35	12	60	SY20
SakSTAR(E65T,K74R,E80A,D82A,K130T,K135R)	30	94	7.2	9.3	9.9	5.6	32	22	15	8.2	11	2.0	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.0	58	24	69	SY21
SakSTAR(S34G,G36R,K74R,K130T,K135R)		250	5.6	54	2.9	38	25	36	6.5	18	5.5	1.9	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.9	75	33	68	SY10
SakSTAR(E65A,K74R,E80A,D82A,K130T,K135R)	40	140	8.2	10	13	5.7	55	24	22	11	15	2.2	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.0	51	17	66	SY18
SakSTAR(E65N,K74R,E80A,D82A,K130T,K135R)	88	120	8.5	12	11	7.0	36	43	13	12	7.9	2.4	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.9	60	29	67	SY23
SakSTAR(E65Q,K74R,E80A,D82A,K130T,K135R)	55	140	9.0	16	16	9.9	59	18	4.5	4.3	10	2.5	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.8	54	22	63	SY22
SakSTAR(K57A,E58A,E61A,E80A,D82A,K130T,K135R)	24	110	2.4	17	2.9	13	12	16	7.8	16	4.5	1.6	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.7	75	64	62	SY13
SakSTAR(E65A,A72S,K74R,E80A,D82A,K130T,K135R)	92	62	8.2	23	3.8	8.4	19	30	11	13	3.6	3.1	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.2	51	13	66	SY53
SakSTAR(E65D,K74Q,E80A,D82A,K130T,K135R)	84	110	7.0	3.9	2.6	4.1	7.2	29	28	14	6.8	2.2	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	0.9	43	13	64	SY30
SakSTAR(E65Q,K74Q,E80A,D82A,K130T,K135R)	54	120	5.1	16	3.2	14	3.7	16	20	2.4	5.7	1.7	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.0	43	21	64	SY47
SakSTAR(K35A,E65D,K74Q,E80A,D82A,K130T,K135R)	56	140	5.1	6.8	2.6	9.5	1.7	28	17	14	8.4	2.1	<0.1	<0.1	<0.1	<0.1	<0.1	<0.1	1.0	35	8	58	SY46

Table 7 - cont'd: Combination mutants of SakSTAR(E80A,D82A,K130T,K135R) with K35A, G36R, E65X, K74X, and selected other amino acids

Variant	Exp. (mg/mL)	Spec. Act. (kU/mg)	murine MAbs															SakSTAR patient plasma					Code
			Epitope cluster I					Epitope cluster II					Epitope cluster III					Pool 10	Subpool B	Subpool C	Pool 40		
			17G11	26A2	30A2	2B12	3G10	18F12	14H5	28H4	32B2	7F10	7H11	25E1	40C8	24C4	1A10						
SakSTAR(K74R,E80A,D82A,S103A,K130T,K135R)	32	160	4.9	22	5.8	14	4.3	26	6.9	7.4	5.0	1.7	<0.1	<0.1	<0.1	<0.1	67	32	69	70	SY24		
SakSTAR(K35A,E65D,K74R,E80A,D82A,E108A,K109A,K130T,K135R)	9.0	89	5.8	30	2.6	26	10	13	16	13	3.2	1.8	<0.1	<0.1	<0.1	<0.1	55	10	63	47	SY12		
SakSTAR(K35A,E65D,K74R,E80A,D82A,E108A,K130T,K135R)	20	91	6.4	20	5.0	15	3.9	22	17	7.4	2.4	1.9	<0.1	<0.1	<0.1	<0.1	44	8	70	53	SY32		
SakSTAR(E65D,K74R,E80A,D82A,E108A,K130T,K135R)	4	90	8.1	6.7	6.9	4.3	29	19	31	11	14	2.1	<0.1	<0.1	<0.1	<0.1	52	11	69	-	SY33		
SakSTAR(K35A,E65D,K74R,E80A,D82A,K109A,K130T,K135R)	42	84	5.5	18	5.3	14	1.6	18	12	7.7	10	1.7	<0.1	<0.1	<0.1	<0.1	43	6	61	50	SY36		
SakSTAR(E65D,K74R,E80A,D82A,K109A,K130T,K135R)	60	130	9.7	6.6	6.8	4.2	28	11	32	12	17	2.3	<0.1	<0.1	<0.1	<0.1	56	10	64	53	SY37		
SakSTAR(K35A,E65D,K74R,E80A,D82A,K130T,K135R,K136A)	28	81	4.5	12	3.3	11	1.7	22	13	7.6	4.9	1.6	<0.1	<0.1	<0.1	<0.1	40	14	52	40	SY21		
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R,K136A)	60	100	6.8	5.8	4.4	4.5	15	33	32	14	7.9	2.0	<0.1	<0.1	<0.1	<0.1	46	28	67	45	SY38		
SakSTAR(E65Q,K74Q,D82A,S84A,K130T,K135R)	170	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	45	21	60	45	SY50N		
SakSTAR(K35A,E65D,K74R,E80A,D82A,K86A,K130T,K135T)	68	86	4.4	20	5.5	15	1.5	15	12	6.4	6.7	1.9	<0.1	<0.1	<0.1	<0.1	36	7	60	55	SY40		
SakSTAR(K35A,K74Q,E80A,D82A,K130T,K135R)	72	120	6.1	3.4	2.5	3.0	5.9	38	14	9.8	6.8	1.9	<0.1	<0.1	<0.1	<0.1	49	16	64	48	SY28		
SakSTAR(K35A,E65D,K74R,E80A,D82A,K130T,K135R)	54	190	8.1	7.5	6.9	5.5	25	37	34	14	7.7	2.3	<0.1	<0.1	<0.1	<0.1	56	28	68	55	SY29		
SakSTAR(K35A,E65D,K74R,E80A,D82A,V132R,K135R)	13	55	6.7	23	5.3	17	2.3	47	19	19	5.1	2.0	<0.1	<0.1	<0.1	<0.1	53	20	88	62	SY61		
SakSTAR(K35A,E65D,K74R,E80A,D82A,T129A,K135R)	13	61	7.0	13	5.1	31	12	27	12	11	6.7	2.5	<0.1	<0.1	<0.1	<0.1	56	18	79	60	SY62		
SakSTAR(K35A,E65D,K74R,E80A,D82A,T129A,K135A)	23	21	6.9	27	5.8	32	20	29	6.6	9.7	5.4	2.1	<0.1	<0.1	<0.1	<0.1	56	17	91	60	SY64		

Association constants ≥ 10 -fold lower and antibody absorption ≤ 60 percent of wild-type SakSTAR are represented in bold type; $\geq 100,000$ HU/mg represented in bold type. NT: not tested.

Table 8: SakSTAR variants with intact specific activity (≥ 100 kHU/mg) and ≤ 50 percent absorption of human antibodies elicited by treatment with wild-type SakSTAR

Variant	Spec. Act. (kU/mg)	SakSTAR patient plasma			
		Pool 10	Subpool B	Subpool C	Pool 40
					Code
SakSTAR(K74Q,K130T,K135R)	190	50	25	67	62
					SY41
SakSTAR(E65A,K74Q,K130T,K135R)	170	45	16	77	55
					SY48
SakSTAR(E65Q,T71S,K74Q,K130T,K135R)	210	49	21	64	59
					SY65
SakSTAR(E65Q,K74Q,E118A,K130A,K135R)	180	50	28	72	58
					SY73
SakSTAR(E65Q,K74Q,N95A,E118A,K130A,K135R)	190	48	27	74	58
					SY74
SakSTAR(K35A,E65Q,K74Q,K130A,K135R)	110	49	26	63	45
					SY75
SakSTAR(E65Q,K74Q,K109A,K130T,K135R)	210	50	22	68	51
					SY81
SakSTAR(K74Q,E80A,D82A,K130T,K135R)	110	46	17	60	48
					SY15
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)	140	43	11	68	57
					SY19
SakSTAR(E65S,K74R,E80A,D82A,K130T,K135R)	110	35	12	60	-
					SY20
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R,K136A)	100	46	28	67	45
					SY35
SakSTAR(K35A,K74Q,E80A,D82A,K130T,K135R)	120	49	16	64	48
					SY28
SakSTAR(E65D,K74Q,E80A,D82A,K130T,K135R)	110	43	13	64	42
					SY30
SakSTAR(E65Q,K74Q,E80A,D82A,K130T,K135R)	120	43	21	64	42
					SY47
SakSTAR(E65Q,K74Q,D82A,S84A,K130T,K135R)	170	45	21	60	45
					SY50N
SakSTAR(K35A,E65D,K74Q,E80A,D82A,K130T,K135R)	140	35	8	58	40
					SY46
SakSTAR(K74Q,K130T,K135R)	190	50	25	67	62
					SY41
SakSTAR(T21A,K35A,E65Q,K74Q,K130A,K135R)	110	50	26	72	50
					SY78
SakSTAR(E65Q,K74Q,K109A,K121A,K130A,K135R)	140	50	31	73	52
					SY88
SakSTAR(E65Q,K74Q,D82A,S84A,K109A,K130A,K135R)	180	43	20	62	44
					SY89

Table 8 - cont'd: SakSTAR variants with intact specific activity (≥ 100 kHU/mg) and ≤ 50 percent absorption of human antibodies elicited by treatment with wild-type SakSTAR

Variant	Spec. Act. (kU/mg)	SakSTAR patient plasma				Code
		Pool 10	Subpool B	Subpool C	Pool 40	
SakSTAR(E65Q,K74Q,N95A,E118A,K130A,K135R,+137A)	120	45	30	74	60	SY93
SakSTAR(E65Q,K74Q,N95A,E118A,K130A,K135R,K136A,+137K)	1,400	37	16	70	54	SY94
SakSTAR(E65Q,K74Q,D82A,S84A,E108A,K109A,K130A,K135R)	110	46	26	63	41	SY95

Antibody absorption ≤ 60 percent of wild-type SakSTAR are represented in bold type; $\geq 100,000$ HU/mg represented in bold type.

Table 2: Fibrinolytic properties of selected SakSTAR variants in human plasma in vitro

Compound	Fibrinolytic potency (C50 in $\mu\text{g/mL}$)	Residual fibrinogen at C50 (% of baseline)	Fibrinogenolytic potency (C50 in $\mu\text{g/mL}$)
SakSTAR	0.18 ± 0.01	93 ± 3.5	24 ± 3.6
SakSTAR(K74Q,E80A,D82A,K130T,K135R)	0.15 ± 0.01	97 ± 3.0	14 ± 3.2
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)	0.24 ± 0.04	94 ± 10	29 ± 3.1
SakSTAR(K35A,E65D,K74Q,E80A,D82A,K130T,K135R)	0.11 ± 0.01	92 ± 3.0	20 ± 2.0
SakSTAR(E65Q,K74Q,N95A,E118A,K130A,K135R,K136A,+137K)	0.13	91	

The data represent mean \pm SD of 3 experiments.

C₅₀: amount of wild type or variant SakSTAR required for 50% clot lysis or 50% fibrinogen breakdown in 2 hrs.

Table 10: Pharmacokinetic parameters of the disposition of staphylokinase-related antigen from plasma following bolus injection of SakSTAR variants (100 µg/kg) in hamsters.

Variant	C ₀ (µg/mL)	A (µg/mL)	B (µg/mL)	t _{1/2} (α) (min)	t _{1/2} (β) (min)	V _C (mL)	AUC (µg.min.mL ⁻¹)	Cl _p (mL.min ⁻¹)
SakSTAR	0.8 ± 0.1	0.6 ± 0.1	0.2 ± 0.0	2.8	7.0	13 ± 1.0	4.6 ± 0.4	2.2 ± 0.2
SakSTAR(K74Q,E80A,D82A,K130T,K135R)	0.5 ± 0.1	0.4 ± 0.1	0.1 ± 0.0	2.0	10	20 ± 2.2	2.5 ± 0.3	4.1 ± 0.5
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)	0.6 ± 0.0	0.5 ± 0.0	0.1 ± 0.0	2.0	10	16 ± 1.1	2.8 ± 0.2	3.7 ± 0.3
SakSTAR(K35A,E65DK74Q,E80A,D82A,K130T,K135R)	1.1 ± 0.1	1.0 ± 0.1	0.1 ± 0.0	2.0	24	9.6 ± 0.7	6.4 ± 0.5	1.6 ± 0.1

Data are mean ± SEM of 4 experiments.

Table 11: Baseline characteristics and treatment outcome of the patients with peripheral arterial occlusion treated with SakSTAR, SakSTAR(K74Q,E80A,D82A,K130T,K135R) or SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)

Compound Patient Id.	Gender	Age (yrs)	Clinical ischemia	Locus of occlusion	Age of occlusion (days)	Length of occlusion (cm)	Recanalization by thrombolysis	Total dose of thrombolytic agent (mg)	Total duration of infusion (hrs)	Additional therapy
SakSTAR										
PUT	M	66	Subacute	Femoro-femoral graft	6	5	Complete	2	23	Stenting left IF artery
VERM	M	73	Acute	Right PA	2	6	Partial	13	23	Right upper leg amputation
GEIV	V	63	Restrain	Left SFA	10	5	Complete	8	7	PTA
POL	M	46	Subacute	Right SFA	30	50	Partial	22	29	Lumbar sympathectomie
BUE	F	53	Claudication	Right AF graft	1	15	Complete	10	13	Desobstruction
VUJ	F	75	Subacute	Left FT graft	2	34	Complete	7	10	PTA
REN	M	48	Restrain	Right IF graft	4	20	Complete	6.5	5	Left AF graft
COR	V	78	Acute	Left AFS	14	9	Complete	4	3	PTA
MAN	M	67	Restrain	Left tibial artery	1	6	Partial	6	5	-
STRA	M	66	Claudication	Right FP graft	14	16	Complete	19	26	-
VANH	M	38	Acute	Left radial artery	4	1	Complete	6	5	-
VANW	F	57	Acute	Right FP graft	1	25	Complete	20	24	New right FP graft
BRA	M	57	Acute	Left FT graft	1	30	Complete	25	43	-
DON	M	60	Claudication	Left FT graft	1	20	Complete	13	19	PTA + stenting
CAM	M	77	Restrain	Right SFA graft	8	30	Complete	27	44	FF graft
Mean ± SEM		62 ± 3.1			6.6 ± 2.1	18 ± 3.5		13 ± 2.1	19 ± 3.5	
SakSTAR(K74Q,E80A,D82A,K130T,K135R)										
IMB	M	66	Claudication	Left SFA	30	5	Complete	24	24	PTA
AZY	M	44	Subacute	Right C.I.A.	7	8	Complete	18	23	Stenting
VIN	M	51	Acute	Right E.I.A.	5	70	Complete	24	30	-
STRO	M	53	Claudication	Left FP junction	14	5	Partial	3.5	2	Aspiration thrombectomy, PTA
VERG	M	62	Restrain	Left SFA	20	6	Complete	19	25	FP bypass
GIE	M	76	Acute	Right FP bypass	2	15	Complete	8.5	17	-
Mean ± SEM		59 ± 4.7			13 ± 4.3	18 ± 10		16 ± 3.4	20 ± 4.0	
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)										
URB	M	57	Subacute	Right E.I.A.	4	8	Complete	8	6	Pseudo aneurysm, right AF graft revision
COM	M	59	Restrain	Right AF graft	7	65	Complete	16	22	-
HAC	M	70	Restrain	Left anterior tibial artery	7	15	Complete	12	14	-
DEW	F	76	Restrain	SFA	21	6	Complete	6	4	-
VAI	F	65	Subacute	Left PA	25	10	Partial	8	6	Aspiration thrombectomy
FIL	M	76	Claudication	Right SFA	28	8	Complete	24	31	PTA
Mean ± SEM		67 ± 3.4			15 ± 4.3	19 ± 9.4		12 ± 2.8	14 ± 4.4	

AF: aortofemoral; CABG: coronary artery bypass graft; CAD: coronary artery disease; CIA: common iliac artery; COPD: chronic obstructive pulmonary disease; DM: diabetes mellitus; EIA: external iliac artery; FF: femorofibular; FP: femoropopliteal; FT: femorotibial; IA: iliac artery; IF: iliofemoral; occi: occlusion; PA: popliteal artery; PTA: percutaneous transluminal angioplasty; SFA: superficial femoral artery; TA: tibial artery; TF: tibiofibular; SC: subclavian.

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Treatment	Absorbant	Insolubilized compound		
		SakSTAR	SakSTAR(K74Q,E80A,D82A,K130T,K135R)	SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R)
SakSTAR (Pool 40)				
SakSTAR		95		
SakSTAR(K74Q,E80A,D82A,K130T,K135R)		48		
SalSTAR(E65D,K74R,E80A,D82A,K130T,K135R)		57		
SakSTAR(K74Q,E80A,D82A,K130T,K135R) (Imb., Vin., Ver., Gie.)				
SakSTAR		94	95	95
SakSTAR(K74Q,E80A,D82A,K130T,K135R)		91	93	89
SalSTAR(E65D,K74R,E80A,D82A,K130T,K135R)		92	94	94
SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) (Urb.)				
SakSTAR		90	88	85
SakSTAR(K74Q,E80A,D82A,K130T,K135R)		94	95	94
SalSTAR(E65D,K74R,E80A,D82A,K130T,K135R)		94	95	94

Data represent median values of the percent absorption with 250 nM absorbant, measured by residual binding to insolubilized compound.

* p= versus SakSTAR; ** p= versus SakSTAR(K74Q,E80A,D82A,K130T,K135R)); and p= versus SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) by paired non-parametric test.

ization level (%)	PEG derivatization
0	none
0	none
0	1
0	none
>60	none
>95	none

APPENDIX

Table 14: Cysteine substitution variants of SakSTAR

Variant	Spec. Act. (kU/mg)	SakSTAR patient plasma				Code
		Pool 10	Subpool B	Subpool C	Pool 40	
SakSTAR(K96C)	790	95	95	95		
SakSTAR(K102C)	280	95	95	95		
SakSTAR(K109C)	1,900	95	95	94	90	
SakSTAR(K35C)						
SakSTAR(K74C)						
SakSTAR(.....)						
SakSTAR(K35A,E65D,K74Q,E80A,D82A,K109C,K130T,K135R)	600	46	32	56	36	SY100
SakSTAR(E65Q,K74Q,D82A,S84A,K109C,K130T,K135R)	1,100	54	53	77	41	SY102

Antibody absorption ≤60 percent of wild-type SakSTAR are represented in bold type; ≥ 100,000 HU/mg represented in bold type.

CLAIMS

1. Staphylokinase derivatives showing a reduced immunogenicity as compared to wild-type staphylokinase, after administration to patients with arterial thrombosis.
2. Staphylokinase derivatives as claimed in claim 1 having essentially the amino acid sequence as depicted in figure 1 in which one or more amino acids have been replaced by another amino acid thus reducing the reactivity with a panel of murine monoclonal antibodies.
3. Staphylokinase derivatives as claimed in claim 1 having essentially the amino acid sequence as depicted in figure 1 in which one or more amino acids have been replaced by another amino acid thus reducing the absorption of SakSTAR-specific antibodies from plasma of patients treated with staphylokinase.
4. Staphylokinase derivatives as claimed in claim 1 having essentially the amino acid sequence as depicted in figure 1 in which one or more amino acids have been replaced by other amino acids, without reducing the specific activity by more than 50 percent.
5. Staphylokinase derivatives SakSTAR(K35X,G36X,E65X,K74X,E80X,D82X,K102X,E108X,K109X,K121X,K130X, K135X,K136X,+137X) having the amino acid sequence as depicted in figure 1 in which the amino acids Lys in position 35, Gly in position 36, Glu in position 65, Lys in position 74, Glu in position 80, Asp in position 82, Lys in position 102, Glu in position 108, Lys in position 109, Lys in position 121, Lys in position 130, Lys in position 135 and/or Lys in position 136 have been replaced with other amino acids and/or in which one amino acid has been added at the COOH-terminus, thus altering the immunogenicity after administration in patients, without markedly reducing the specific activity.
6. Staphylokinase derivatives listed in Tables 1,3,4,5,6,7 and 8, having the amino acid sequence as depicted in figure 1 in which the indicated amino acids have been replaced by

other amino acids thus reducing the absorption of SakSTAR-specific antibodies from plasma of patients treated with staphylokinase, without reducing the specific activity.

7. Staphylokinase derivatives having an amino acid substituted with Cys, resulting in dimerization, increased specific activity, delayed clearance and/or increased thrombolytic potency.
8. Staphylokinase derivatives with polyethylene glycol substitution....
9. Method for producing the staphylokinase derivatives as claimed in claims 1 to 7, comprising the steps of:
 - a. preparing a DNA fragment comprising at least the part of the coding sequence of staphylokinase that provides for its biological activity;
 - b. performing in vitro site-directed mutagenesis on the DNA fragment to replace one or more codons for wild-type amino acids by a codon for another amino acid;
 - c. cloning the mutated DNA fragment in a suitable vector;
 - d. transforming or transfecting a suitable host cell with the vector; and
 - e. culturing the host cell under conditions suitable for expressing the DNA fragment.
10. Method as claimed in claim 9, wherein the DNA fragment is a 453 bp *EcoRI-HindIII* fragment of the plasmid *pMEX602sakB*, the *in vitro* site-directed mutagenesis is performed and the mutated DNA fragment is expressed in *E. coli*.
11. Pharmaceutical composition comprising at least one of the staphylokinase derivatives as claimed in claims 1 to 8 together with a suitable excipient.
12. Pharmaceutical composition as claimed in claim 11 for treating arterial thrombosis.

LEGENDS TO THE FIGURES

- Fig 1. Protein sequence of wild-type staphylokinase, SakSTAR. Numbering starts with the NH₂-terminal amino acid of mature full length staphylokinase.
- Fig 2. Time course of neutralizing activities (left panel) and specific IgG against administered agent (right panel) following intra-arterial infusion of SakSTAR (open circles, n= 9), SakSTAR(K74A) (closed circles, n= 11) or SakSTAR(K74A,E75A, R77A) (open squares, n= 6) in patients with peripheral arterial occlusion. The data represent median values and interquartile ranges, in µg/mL.
- Fig 3. Protein sequence of wild-type staphylokinase, SakSTAR with indicated amino acid substitutions.
- squares: single amino acid substitutions; circles: combined (2 to 3) amino acid to Ala substitutions.
- Fig. 4. Temperature stability of SakSTAR, (A); SakSTAR(K74Q,E80A,D82A,K130T, K135R), (B); SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R), (C); and SakSTAR(K35A,E65D,K74Q,E80A,D82A, K130T,K135R), (D).
- (○): 4°C; (●): 20°C; (▽): 37°C; (▼): 56°C; (□): 70°C.
- Fig 5. Time course of neutralizing activities (left panel) and specific IgG against administered agent (right panel) following intra-arterial infusion of SakSTAR (circles, n=), SakSTAR(K74Q,E80A,D82A,K130T,K135R) (squares, n= 6) or SakSTAR(E65D,K74R,E80A,D82A,K130T,K135R) (triangles, n= 6) in patients with peripheral arterial occlusion. The data represent median values and 15-85 percentile ranges, in µg/mL.

Methods for the identification, production and use of staphylokinase derivatives characterized by a reduced immunogenicity after administration in patients. The derivatives of the invention are obtained by preparing a DNA fragment comprising at least the part of the coding sequence of staphylokinase that provides for its biological activity; performing *in vitro* site-directed mutagenesis on the DNA fragment to replace one or more codons for wild-type amino acids by a codon for another amino acid; cloning the mutated DNA fragment in a suitable vector; transforming or transfecting a suitable host cell with the vector; culturing the host cell under conditions suitable for expressing the DNA fragment; and purifying the expressed staphylokinase derivative to homogeneity. Preferably the DNA fragment is a 453 bp *EcoRI-HindIII* fragment of the plasmid *pMEX602sakB*, (*pMEX.SakSTAR*), the *in vitro* site-directed mutagenesis is performed by spliced overlap extension polymerase chain reaction and the mutated DNA fragment is expressed in *E. coli* strain TG1 or WK6. The invention also relates to pharmaceutical compositions comprising at least one of the staphylokinase derivatives according to the invention together with a suitable excipient, for treatment of arterial thrombosis.